

The metabolic cost of egg production is repeatable

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Summary

The metabolic cost of egg production in birds (passerines) has been measured as a 16–27% increase in basal or resting metabolic rate (BMR and RMR, respectively) when comparing non-breeding values with those in egg-producing individuals. However, available data to date have been obtained in free-living birds and may thus be confounded by the effect of variable ecological conditions on non-reproductive physiological machinery (organ mass or metabolic activity) which might contribute to measured variation in RMR. Here, we show that in captive, controlled conditions, the process of egg formation induces a 22% increase in RMR in female zebra finches *Taeniopygia guttata*. Among individuals, variation

in laying RMR is independent of egg mass, clutch size or total clutch mass. Importantly, we show that individual variation in both non-breeding and laying RMR is repeatable over periods of at least 8–10 months, i.e. individual variation in RMR remained constant over time for any given physiological state. This suggests that the metabolic cost of egg formation should respond to selection. However, we also show that in males, but not females, repeatability of RMR declines over time even when birds are kept in constant controlled conditions.

Key words: basal metabolic rate, resting metabolic rate, repeatability, egg production, reproduction, zebra finch, *Taeniopygia guttata*.

Introduction

Basal metabolic rate (BMR) is a measure of the overall energy consumed by the complete array of organs and physiological systems comprising an animal in resting, post-absorptive conditions at thermoneutrality. Although often considered to represent a minimal level of metabolism, BMR is, in fact, a highly flexible phenotypic trait and fluctuates in relation to adjustments in internal physiological machinery (Piersma, 2002), such as when body composition varies with physiological state (e.g. reproduction and migration; Battley et al., 2000, 2001; Silverin, 1981; Vézina and Williams, 2003) and/or changing ecological conditions (Drobney, 1984; Geluso and Hayes, 1999; Heitmeyer, 1988; Rogers et al., 1993; Hilton et al., 2000; Hammond et al., 2001). Although many studies have now documented sources of variation in BMR, few studies have focused on individual variation or repeatability of BMR either for free-living or captive animals (Bech et al., 1999; Horak et al., 2002; Labocha et al., 2004). Individual variation provides the raw material on which selection acts, but for phenotypic traits to evolve through natural selection between-individual variation must be consistent, i.e. it must show high and significant repeatability. Repeatability is therefore a very useful tool for quantifying the extent to which an individual's behaviour or physiology remains consistent over time and can be informative in determining how effective selection will be in changing traits over time (Bennett, 1987; Falconer and Mackay, 1996; Dohm, 2002).

Here, we show that resting metabolic rate (RMR; our estimator of BMR, see Materials and methods) is repeatable in relation to physiological state, in non-breeding and egg-producing zebra finches *Taeniopygia guttata*. We have previously shown, in European starlings *Sturnus vulgaris*, that egg formation induces a 22% increase in RMR over pre-reproductive values (Vézina and Williams, 2002), which is comparable to data reported for other free-living avian species (house sparrows *Passer domesticus*, 16% over non-breeding BMR; Chappell et al., 1999; great tits *Parus major*, 27% over-wintering RMR; Nilsson and Raberg, 2001). However, these observations were made on free-living animals where RMR might be confounded by variation in ecological conditions *via* effects on non-reproductive physiological 'machinery' (organ mass or metabolic activity), which might independently contribute to measured variation in RMR (Vézina and Williams, 2003). We conducted a study in controlled, laboratory conditions, allowing for an unbiased estimate of the metabolic cost of egg production, eliminating the confounding effects of natural variations in ecological condition. We predicted that there would be a similar state-dependent change in RMR in laying zebra finches allowing us to (1) estimate the energetic investment associated with reproductive effort measured as egg production in zebra finches, (2) determine relationships between RMR in laying birds and measures of reproductive output (egg and clutch size), (3) estimate

repeatability for non-breeding and laying RMR to evaluate the level of stability of this trait within individuals and between breeding attempts, and (4) investigate the effect of time on repeatability of RMR over short (8 days) and long (10 months) timescales.

Materials and methods

Animal husbandry

Zebra finches (*Taeniopygia guttata*, Vieillot) were kept in controlled environmental conditions (temperature 19°–23°C; humidity 35%–55%; constant light intensity, 14 h:10 h L:D, with lights on at 0800 h). All birds were maintained on a mixed-seed diet (Panicum and white millet, 50:50; approximately 12.0% protein, 4.7% lipid; Jamieson's Pet Food, Vancouver, BC, Canada) with water, grit and cuttlefish bone *ad libitum*, and were given a multivitamin supplement in the drinking water once per week. All non-breeding birds were kept with the same sex in 61×46×41 cm cages. Birds forming breeding pairs were chosen randomly and were housed in the same type of cage and provided with an external nest box (11.5 cm×11.5 cm×11.5 cm). Nest boxes were checked daily between 1000–1200 h, and all new eggs were weighed (to ±0.001 g) and numbered. A clutch was considered complete after two consecutive days with no new eggs. All breeding pairs had access to an egg food supplement replaced daily (20.3% protein, 6.6% lipid). All experiments and animal husbandry were carried out under a Simon Fraser University Animal Care Committee permit (692B-94), following the guidelines of the Canadian Committee on Animal Care.

Measurement of resting metabolic rate

Basal metabolic rate is defined as the energy consumed by a resting post-absorptive animal during the inactive phase of the circadian cycle at a temperature within the thermoneutral range for the animal (Commission for Thermal Physiology of the IUPS, 2001). However, since laying birds are producing eggs, they have to be considered to be in an 'active physiological state' (Vézina and Williams, 2002). We therefore consider the term *resting* metabolic rate more appropriate in the present study. All resting metabolic rate (RMR) measurements were completed using a flow-through respirometry set-up (Sable Systems International; oxygen analyzer model FC-1, CO₂ analyser model CA-1) described elsewhere (Vézina et al., 2003). Birds were taken from their cages within 10–15 min after lights were turned off, their body mass measured (±0.1 g), and were placed randomly into one of four metabolic chambers (1.5 l) for approximately 1 h prior to the beginning of RMR measurements. All chambers continuously received 500 ml min⁻¹ of dry CO₂-free air and were kept in the dark at 35°C, which is within the thermoneutral zone for this species (lower critical temperature=33°C; Meijer et al., 1996). RMR measurements were always started at 23:00 h. All measurement sequences started by recording 20 min of ambient baseline air. After

baseline recording, the out-flowing air from the first chamber was sampled for 33 min before switching to sampling the second chamber for 33 min. Then the system sampled baseline air for 10 min before changing to the third and fourth chambers. This cycle was repeated three times over the night (with ten minutes of baseline in between each set of two chambers), giving 99 min of recording per chamber over 8 h. After RMR measurement, the birds were re-weighed and placed back into their cage (approximately 30 min to 1 h before lights were turned on). To calculate RMR, the average of first and second masses was used and \dot{V}_{O_2} was calculated using a running mean representing 10 min of recording, with the lowest average taken as RMR. This value was always found in the last 5 h of the night. Preliminary analysis showed that measuring RMR using this protocol did not generate a time effect (Hayes et al., 1992; ANOVA testing for chamber position in the measurement sequence on mass-corrected RMR: $F_{3,53}=0.7$, $P=0.5$).

Experimental groups

To evaluate the relative increase in resting metabolic rate associated with egg production in zebra finches, we compared RMR values for a given female measured as non-breeder (NB; maintained in a single sex group), at laying (LY; day of first egg laid) and at 17 days into the chick-rearing period (CK; approximately 4 days before fledging). To estimate repeatability of LY RMR, females were paired again after a resting period and RMR was measured a second time at the one-egg stage. Since we could only measure four birds per night, RMR was recorded for laying birds at the 1-egg ($N=38$) and two-egg ($N=7$) stages to accommodate for days with more than four new layers. RMR did not differ significantly between egg stages (1-egg RMR=52.0±0.9 ml O₂ h⁻¹; 2-egg RMR=50.0±1.7 ml O₂ h⁻¹; $P=0.8$). For repeatability of RMR in non-breeders, birds and cage availability prevented us from measuring RMR repeatedly in all females measured as LY. We, therefore, evaluated repeatability of NB RMR using a group of birds composed of 42% of females that were not used in the breeding protocol. As we were interested in the effect of time on RMR repeatability, in a separate experiment, we measured RMR in a group of non-breeding males twice, with measurements 8 days apart and then a third time 127–249 days later. Mean RMR and body mass are presented in Table 1 with the respective sample sizes for each group.

Statistical analysis

All data were tested to ensure normality (Shapiro-Wilk test; Zar, 1996). To compare within individual changes in RMR from the non-breeding to one-egg and chick rearing stages, we used repeated measures analysis of covariance (ANCOVA) using body mass as a covariate. We used the same method to investigate potential changes in mean RMR measured twice as LY or NB stages. Repeatability of residual RMR (effect of body mass factored out by regression analysis) was calculated following the method proposed by Lessells and Boag (1987).

Table 1. Mean body mass, mean resting metabolic rate and samples size at all breeding stages measured for the four analysis treatments

		Breeding stage*					
		NB ₁	LY ₁	CK	NB ₂	LY ₂	NB ₃
Repeatability males	Body mass (g)	16.1±0.3	–	–	15.4±0.3	–	14.7±0.4
	RMR (ml O ₂ h ⁻¹)	36.1±0.9	–	–	38.4±1.0	–	39.8±0.9
	Sample size	27	–	–	27	–	24
Breeding RMR females	Body mass (g)	15.4±0.2	16.7±0.2	14.1±0.2	–	–	–
	RMR (ml O ₂ h ⁻¹)	39.8±0.6	51.2±0.7	42.3±0.7	–	–	–
	Sample size	57	45	16	–	–	–
Repeatability NB females	Body mass (g)	15.4±0.2	–	–	14.9±0.2	–	–
	RMR (ml O ₂ h ⁻¹)	40.5±0.8	–	–	42.4±0.8	–	–
	Sample size	37	–	–	37	–	–
Repeatability LY females	Body mass (g)	–	17.1±0.4	–	–	17.3±0.4	–
	RMR (ml O ₂ h ⁻¹)	–	52.7±1.5	–	–	52.2±1.4	–
	Sample size	–	19	–	–	19	–

Note: see text and figures for mass-corrected values and analysis of resting metabolic rate (RMR).

*Stages: NB, non breeding bird; LY, day of first egg laid; CK, chick rearing; subscript numbers indicate first, second or third measurement. Values are means ± S.E.M.

Therefore, our repeatability index reflects the amount of variation in RMR among rather than within individuals. *Post-hoc* multiple comparisons between groups were performed using the Bonferroni procedure to reduce the risk of committing type I errors (Rice, 1989). Data are reported as mean ± S.E.M.

Results

Variation in RMR in relation to breeding stage

Repeated measures ANCOVA showed that resting metabolic rate differed significantly between breeding stages when controlling for the effect of body mass ($F_{2,84}=71.0$, $P<0.0001$, no significant interaction between stage and body mass; Fig. 1). Least square mean comparison (controlling for the mass effect) revealed that LY RMR was, on average, 22.4% higher than NB values (*post-hoc* multiple comparison $P<0.0001$) and 8.2% higher than CK values (*post-hoc* multiple comparison $P<0.01$; Fig. 1). Using a larger sample size based on all females for which we have at least one LY RMR measurement combined with egg data ($N=39$), we found no significant relationship between residual LY RMR (correcting the effect of body mass) and mean egg mass, total clutch mass or clutch size ($P\geq 0.4$ in all cases, overall mean egg mass=1.08±0.02 g, clutch mass=6.16±0.2 g and clutch size=6±0.2 eggs).

Repeatability of RMR

Our data on non-breeding RMR in male zebra finches allows for the investigation of short- and long-term repeatability. Comparing residual (mass corrected) RMR for the first and second measurements over an 8 day period, 62.6% of the total variation was explained by among-

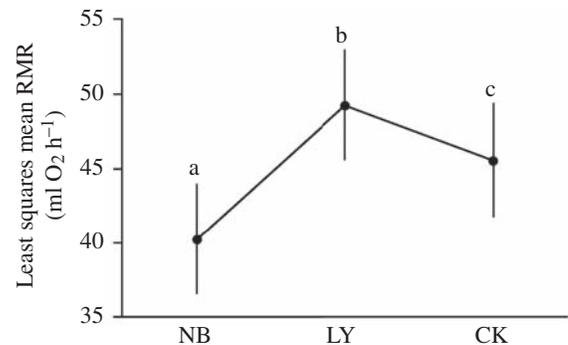


Fig. 1. Least square mean resting metabolic rate (RMR) in non-breeding (NB), laying (LY) and chick-rearing (CK) stage of female zebra finches. Values are computed from a repeated measures ANCOVA controlling for the effect of body mass. Different letters indicate significant differences between groups.

individual variability ($F_{26,27}=4.4$, $P<0.0001$; Fig. 2A). However, repeatability of RMR declined with time (measure 2 vs 3; 127 to 249 days; 44.5%; $F_{23,24}=2.6$ $P<0.05$; Fig. 2B, measure 1 vs 3; 135 to 257 days; 28.7%; $F_{23,24}=1.8$, $P=0.08$; Fig. 2C).

In non-breeding females, we measured RMR twice over a period of time spanning 13–311 days. Repeatability of residual RMR in these birds was 51.6% ($F_{36,37}=3.1$, $P<0.0005$; Fig. 3A). Including the delay between measurements in a multiple regression model showed that the relationship between first and second RMR measurements was still significant when time was taken into account (time effect $P<0.05$, RMR1 effect on RMR2: $P<0.001$, no significant interaction term; overall model $r^2=0.35$, $N=37$, $P<0.001$). Repeated measures ANCOVA showed a 5.9% increase

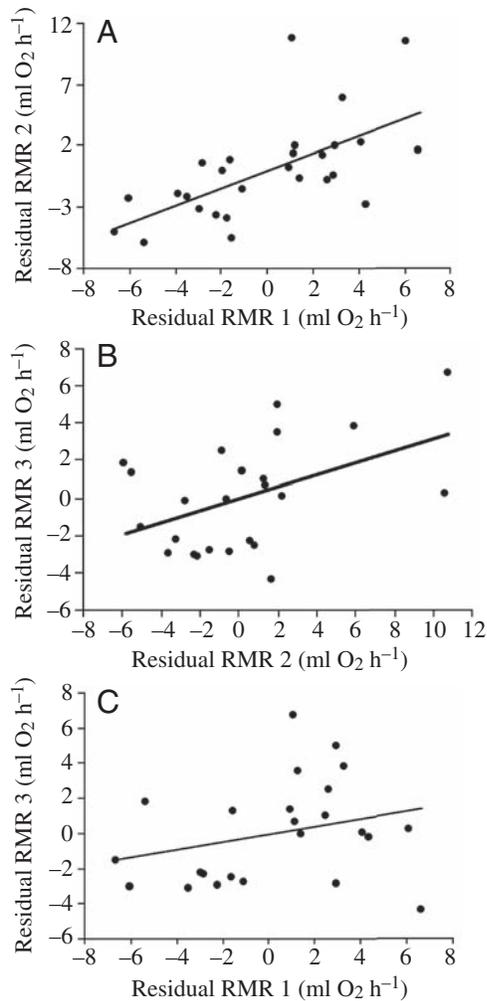


Fig. 2. Correlation between residual resting metabolic rate (RMR) measured repeatedly in male zebra finches. (A) First vs second measurement, (B) second vs third measurement, (C) first vs third measurement. Residuals are controlling for the effect of body mass.

between first and second RMR measurement when controlling for body mass ($F_{1,37}=10.3$, $P<0.005$; Fig. 4). Comparing data on initial NB RMR for males and females, we found that RMR was independent of sex when controlling for body mass (ANCOVA; $P=0.3$, no significant interaction term between sex and body mass).

We measured RMR in laying females twice over a period spanning 38 to 254 days. Repeatability of residual RMR was 52.6% ($F_{18,19}=3.22$, $P<0.01$; Fig. 3B) and multiple regression showed no effect of delay between measurements in this particular data set (time effect $P=0.8$). Repeated measures ANCOVA showed no significant differences between the first and second LY RMR measurement when controlling for the effect of body mass ($P=0.4$, no significant interaction between body mass and measurement sequence; Fig. 4) meaning that absolute mean LY RMR values did not change between breeding attempts.

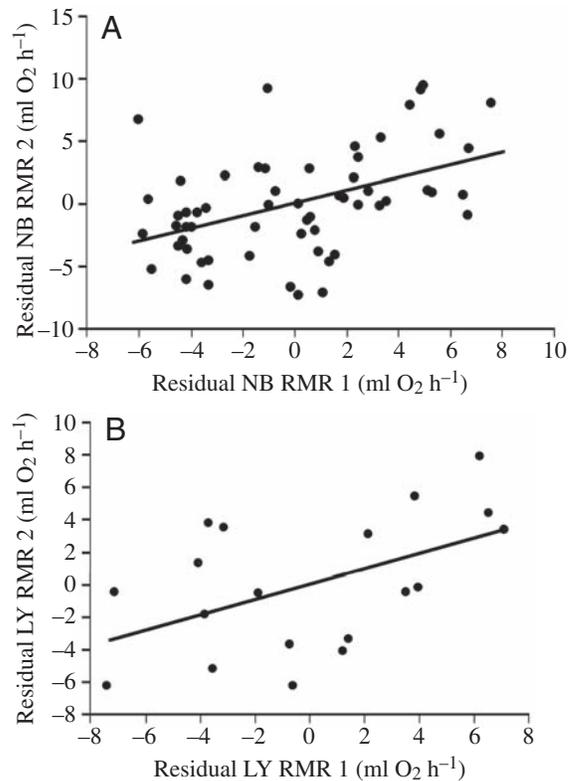


Fig. 3. Correlation between first and second resting metabolic rate (RMR) measured at (A) the non-breeding (NB) stage and (B) the laying (LY) stage in female zebra finches. Residuals correct for the effect of body mass.

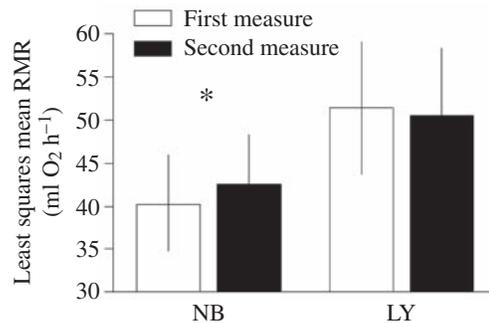


Fig. 4. Least square mean resting metabolic rate (RMR) measured twice in non-breeding (NB) and laying (LY) females zebra finches. Values are computed from a repeated measures ANCOVA and control for the effect of body mass. An asterisk indicates significant difference between measurements within stage.

Discussion

Our study clearly demonstrates that for female zebra finches living in controlled environmental conditions with food *ad libitum*, the physiological changes associated with egg production induce a significant (22%) increase in resting metabolic rate over non-reproductive value, an increase comparable to earlier reports (Chappell et al., 1999; Nilsson and Raberg, 2001; Vézina and Williams, 2002). Although

previous studies involved wild birds subjected to variations in local ecological conditions, potentially impacting non-reproductive physiology (Vézina and Williams, 2002, 2003), the reported increases in metabolic rate associated with egg production in previous studies are very similar to our measurements in captive birds. This suggests that the effect of egg-production on metabolic rate, reported here and elsewhere, truly reflects physiological changes associated with the development of reproductive physiological systems, and is relatively independent of ecological conditions. However, the exact mechanism responsible for the elevated energy consumption has not yet been clearly identified. The mass of female reproductive organs was found to be positively related to reproductive BMR in house sparrows ($r^2=0.30$; figure 3 in Chappell et al., 1999). Similarly, lean dry oviduct mass explained 18% of the variation in elevated mass-corrected laying RMR in starlings (Vézina and Williams, 2003) and fresh oviduct mass explained 23% of the variation in mass-corrected RMR in pre-laying zebra finches (F.V., unpublished data: $r^2=0.23$, $N=20$, $P<0.05$). Nevertheless, these results suggest that at least 70% of the variation in egg-producing RMR still remains unexplained. Vézina et al. (2003) demonstrated that, in captive zebra finches, producing vitellogenin and yolk-targeted very low density lipoprotein (VTG and VLDL_y, respectively) – the main yolk precursors (Williams, 1998) – does not have a detectable cost in terms of changing RMR. Therefore, other physiological mechanisms must be responsible for part of the increased energy consumption. In the present study, we found no relationship between laying RMR and our reproductive effort parameters: egg mass, clutch size and clutch mass. This result contrasts with the reported positive correlation between overall (not mass-corrected) RMR and mean egg mass in laying great tits (Nilsson and Raberg, 2001) and with the negative relationship between residual RMR and the mass of the oviductal and first egg in European starlings reported in one of three years (Vézina and Williams, 2002). These findings suggest that the size and number of yolky follicles in the ovary at the time of measurement are not consistently related to variation in laying RMR, and clearly more research is needed to fully understand the underlying mechanisms responsible for the metabolic cost of egg production.

An interesting finding is that zebra finches at the chick-rearing stage show a 13% higher RMR compared to non-breeding values. This difference is clearly not the result of partially or non-regressed reproductive organs since this happens very rapidly, within a day of the final ovulation, in this (Williams and Ames, 2004) and other species (Vézina and Williams, 2003). It appears that resting metabolic rate in chick-rearing individuals reflects a physiological state that differs from that in both non-reproductive and egg-producing birds (see also Vézina and Williams, 2003).

Although egg production induces a 22% increase in resting metabolic rate, RMR was repeatable in both non-breeding and laying birds, i.e. individual variation in RMR was consistent for a given physiological state. This may have significant

evolutionary consequences if the increase in RMR associated with egg production is consistently posing a constraint on the animal's energy budget; there is some evidence that this is the case. Williams and Ternan (1999) showed that egg laying zebra finches reduce their level of locomotor activity by 46% while food intake actually decreases by 8%, suggesting that egg production may force females to reallocate energy through behavioural strategies. Furthermore, experiments where females have been induced to lay extra eggs and produce larger clutch sizes showed several fitness consequences visible in lower egg quality affecting chick survival (Monaghan et al., 1995; Nager et al., 2000), lower capacity of females to raise their brood (Heaney and Monaghan, 1995; Monaghan et al., 1998) and lower female survival between years and reproductive success the year following the experiment (Nager et al., 2001; Visser and Lessells, 2001).

In contrast to RMR in egg-producing females, for which repeatability was not affected by time, we found that repeatability of non-breeding RMR decreased with time in both sexes and even that the average value changed between measurements (by 5.9%) in females. Similar declines in repeatability of metabolic rate over time in wild animals have been reported earlier (measurements of BMR: Bech et al., 1999; $\dot{V}_{O_2,max}$: Chappell et al., 1995) but this may not be surprising, given that ecological conditions, and their effects on physiology, may vary between measurements. However, the reason for the increase in interindividual variability in RMR over time in our controlled system is not obvious. There is only one other study that we are aware of that reported time effects on repeatability of BMR in captive birds. Horak et al. (2002) found a decreasing level of repeatability in greenfinches *Carduelis chloris* going from 87% at 8 days between measurements to 63% at 4 months between measurements (table 1 in Horak et al., 2002). In their case, however, the birds were maintained in semi-natural conditions between the two sets of measurements (natural changing photoperiod) which could explain part of the decrease in repeatability. A potential explanation for our finding is that our non-breeding individuals were kept in reserve cages forming large groups (10–20 birds) between measurements. Zebra finches are social birds and will form social hierarchies when maintained in groups (Zann, 1996). It is thus possible that social interactions and access to food when birds are kept in non-breeding same-sex groups, affects some aspects of individual physiology, such as fat content – known for its potential diluting effect on the body mass – metabolic rate relationships (Scott and Evans, 1992) or hormonal state, which may influence behaviour and thus energy expenditure (Ramenofsky, 1984; Wikelski et al., 1999a,b). Therefore, changing group composition between RMR measurements may explain the decrease in RMR repeatability and the change in mean non-breeding values over time, a condition that was not encountered by breeding females. Alternatively, but not exclusively, sex-specific effects of ageing could play a role since mass-specific metabolic rate is known to decline in older individuals (Rolfe and Brown, 1997).

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