Metabolic rate and body size are linked with perception of temporal information

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Body size and metabolic rate both fundamentally constrain how species interact with their environment, and hence ultimately affect their niche. While many mechanisms leading to these constraints have been explored, their effects on the resolution at which temporal information is perceived have been largely overlooked. The visual system acts as a gateway to the dynamic environment and the relative resolution at which organisms are able to acquire and process visual information is likely to restrict their ability to interact with events around them. As both smaller size and higher metabolic rates should facilitate rapid behavioural responses, we hypothesized that these traits would favour perception of temporal change over finer timescales. Using critical flicker fusion frequency, the lowest frequency of flashing at which a flickering light source is perceived as constant, as a measure of the maximum rate of temporal information processing in the visual system, we carried out a phylogenetic comparative analysis of a wide range of vertebrates that supported this hypothesis. Our results have implications for the evolution of signalling systems and predator–prey interactions, and, combined with the strong influence that both body mass and metabolism have on a species’ ecological niche, suggest that time perception may constitute an important and overlooked dimension of niche differentiation.

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All biological systems, from organisms to ecosystems, are shaped by universal constraints. For example, body size and metabolic rate act as important constraints on several characteristics of organisms such as life history and behaviour, making them a particularly common and well-studied aspect of species’ ecology (Brown \textit{et al.} 2004; Woodward \textit{et al.} 2005; Sibly \textit{et al.} 2012). However, constraints imposed by the organism’s sensory limitations are probably equally important and yet frequently overlooked (McGill \& Mittelbach 2006; Pawar \textit{et al.} 2012).

In animal species, the limitations of sensory systems are crucial in shaping both intra- and interspecific interactions. For example, the ability to spot and accurately predict the motion of the opposite party can be pivotal in determining the outcome in both predator–prey interactions (Fig. 1; \textit{Cronin} 2005; \textit{Stevens} 2007; Stevens \textit{et al.} 2011; Clark \textit{et al.} 2012; De Vries \& Clandinin 2012) and the locating of mates (Land \& Collett 1974; Hornstein \textit{et al.} 2000). While the links among sensory limitations, foraging and spatial acuity have been studied in detail (e.g. in the use of search images for prey detection; \textit{Cronin} 2005), the temporal resolution at which dynamic information can be perceived has received considerably less attention, in particular within a general ecological and evolutionary context.

The ability to integrate information over fine timescales, that is, at high temporal resolution, is thus fundamental to many aspects of an organism’s ecology and behaviour. Furthermore, temporal resolution is also directly linked to the perception of the passage of time itself for humans, in particular when tracking fast moving stimuli (\textit{Hagura} \textit{et al.} 2012). From an evolutionary perspective, a trade-off exists between the demand for information at high temporal resolution and the costs of its acquisition given the energetic demands associated with increased rates of neural processing in.
the visual system (Laughlin 2001). This trade-off is likely to be shaped by various ecological (e.g. mode of predation) and environmental factors (e.g. light levels) as well as intrinsic factors (e.g. morphology) that will ultimately shape an organism’s optimal temporal resolution for sensory perception. For example, predators of slow-moving prey may require less temporal resolution than predators that engage in active pursuit of fast-moving prey, such as raptors catching prey during flight.

This ability to perceive and react to a dynamic environment is a key behavioural and ecological trait. Ecologically, interaction strengths can be affected by the ability to identify and track fast-moving objects such as prey or mates (Fig. 1; Land & Collett 1974; Fritsches et al. 2005). The necessity of this ability to perceive one’s environs accurately is perhaps best demonstrated in cases where temporal resolution is too coarse to allow the observer to follow the motion of a moving target accurately. A stark demonstration of this can be seen in the tiger beetle, Cicindela hudsoni, which, owing to the relatively low temporal resolution of its visual system, must take a stop–start approach in order to recalibrate the position of its prey when hunting (Gilbert 1997). In humans, the limitations of our temporal perception are apparent when tracking fast-moving objects such as the curving trajectory of a ball in soccer (Dessing & Craig 2010) and baseball (Bahill & Baldwin 2004).

Two intrinsic factors that may shape the costs and benefits of the temporal resolution of the sensory system, in particular with respect to their effects on an individual’s ability to interact with the environment on short timescales, are body size and metabolic rate. As larger body sizes decrease manoeuvrability (Heglund & Taylor 1988; Dudley 2002; Biewener 2003; Sato et al. 2007; Vogel 2008; Hedrick 2011; Watanabe et al. 2012) and higher metabolic rates increase both manoeuvrability and the physiological ability to process information (Laughlin 2001; Franz & Ronacher 2002), we hypothesized that smaller organisms and those with higher metabolic rates perceive temporal change on finer timescales.

To quantify the temporal perceptual abilities of a range of species we took advantage of the all or nothing nature of neural firing in the visual system. Owing to this binary firing, temporal resolution must be encoded in terms of discrete units, as biological visual systems must discretize the continuous-time and continuous-space information reaching the retina and then integrate this information over some time period. This ‘integration time’ of visual systems can be quantified using the critical flicker fusion frequency (CFF): the lowest frequency of flashing at which a flickering light source is perceived as constant (D’Eath 1998; Schwartz 2009). As light intensity can increase the number of flashes that can be observed per second, the maximum CFF value, as measured in a response curve of CFF against light intensity (Ferry 1892; Porter 1902), can be used as a proxy for the temporal resolution of the sensory system.

We used CFF to compare the temporal resolution of the visual system in a wide range of vertebrate species including representatives from Mammalia, Reptilia, Aves, Amphibia, Elasmobranchii and Actinopterygii. Using phylogenetic comparative methods and controlling for the light levels each species typically experiences, we tested whether the temporal resolution of the sensory system increases with mass-specific metabolic rate and decreases with body mass.

METHODS

Data Collection

To test our prediction that CFF increases with mass-specific metabolic rate and decreases with body size (when controlling for light levels), we collated data on CFF values of vertebrate species from the literature (Table 1). We only included values from studies that measured CFF using either behavioural or electroretinogram (ERG) procedures. In behavioural studies, CFF is measured through conditional training with the subject trained to respond to a change in its perception of a light flashing (D’Eath 1998; Rubene et al. 2010). For example, Lisney et al. (2011) conducted behavioural tests in domestic chickens, Gallus gallus, through choice experiments using flickering and nonflickering stimulus windows with choice of the correct stimulus rewarded with food. This is repeated over a range of light intensities and flicker frequencies until individuals can no longer distinguish between the stimuli. In ERG studies, a direct measurement of the electrical response of the retina in reaction to a flashing light source is used as a measure of CFF (D’Eath 1998; Schwartz 2009). As there may be further processing of temporal information after it reaches the retina that may cause behavioural studies to measure lower CFF values (D’Eath 1998), we included the
Table 1
Data set used in main analysis

| Species                  | CFF  | Mg     | qWg   | Brain mass | Light levels |
|--------------------------|------|--------|-------|------------|--------------|
| Ambystoma tigrinum       | 30.0±2.4 | 10.78±26 | 0.0016±28 | NA          | L            |
| Anguilla anguilla        | 14.8±2.2 | 71.1±28  | 0.0013±28 | NA          | L            |
| Anoto cristatus          | 70.0±3.3 | 6.0±9    | 0.0089±29 | NA          | H            |
| Asio flammeus            | 70.0±4.4 | 406.0±40 | 0.0032±28 | 5.45±9      | H            |
| Bubo virginius           | 45.5±4.5 | 1450.0±31 | 0.0036±28 | 13.7±10     | L            |
| Canis lupus              | 80.0±8.0 | 13900.0±12 | 0.0018±28 | 80.0±7      | H            |
| Carolius auratus         | 67.2±6.7 | 10.8±31  | 0.0013±28 | 0.017±1     | H            |
| Carcharhinus acronotus   | 18.8±8  | 1440.0±8  | 0.0011±46 | NA          | L            |
| Careta caretta           | 40.0±9  | 135000.0±24 | 0.0008±37 | 2.7±90      | H            |
| Cavia porcellus          | 50.0±5   | 620.0±3   | 0.0036±38 | 3.8±12      | L            |
| Chelonias mydas          | 40.0±9  | 128000.0±26 | 0.0025±28 | 8.6±17      | H            |
| Colomba livia            | 100.0±4.4 | 315.0±37 | 0.0045±28 | 2.3±70      | H            |
| Dermotherus coriacea     | 15.0±11 | 354000.0±28 | 0.0043±38 | 30.0±11     | H            |
| Felis catus              | 55.0±12 | 3054.0±32 | 0.0039±40 | 28.4±7      | L            |
| Gallus gallus domesticus | 87.0±13 | 2170.0±39 | 0.0022±24 | 3.6±46      | H            |
| Gekko Gecko              | 20.0±14 | 54.0±9   | 0.0034±28 | 0.27±L      | L            |
| Homo sapiens             | 60.0±15  | 67100.0±41 | 0.0017±30 | 130.0±70    | H            |
| Iguana iguana            | 80.0±10  | 750.0±9   | 0.0029±18 | 0.61±7      | H            |
| Macco Iguana mutata      | 39.0±16 | 771.0±41  | 0.0020±34 | 91±75       | H            |
| Melopsittacus undulates   | 74.0±8.7 | 33.7±28  | 0.0120±48 | 1.5±70      | H            |
| Negaprion brevirostris    | 37.0±18 | 92897.0±44 | 0.0053±28 | NA          | L            |
| Onychocynthus mykiss      | 27.0±19 | 4000.0±45 | 0.0041±28 | 0.5±71      | L            |
| Orizyas latipes           | 37.0±20  | 2.1±9    | 0.0007±28 | 0.017±7     | L            |
| Papagallus ep.GO          | 37.0±12  | 119600.0±46 | 0.0021±31 | 228.5±2     | H            |
| Raja eirinaeae            | 30.0±22  | 500.0±67 | 0.0024±27 | 2.3±71      | L            |
| Rattus norvegicus         | 39.0±23 | 237.0±46  | 0.0076±98 | 2.3±70      | H            |
| Spermochus lateralis      | 120.0±10 | 215.5±49 | 0.0033±54 | 3.6±80      | H            |
| Sphenodon punctatus       | 45.0±24  | 353.0±50 | 0.0017±28 | NA          | L            |
| Sphyra lewini            | 27.0±18  | 1809.0±51 | 0.0016±55 | 60.0±7      | L            |
| Sturnus vulgaris          | 100.0±4.5 | 75.0±28 | 0.012±18 | 1.97±4      | H            |
| Tamias amoenus           | 100.0±10  | 51.9±12  | 0.0037±30 | 1.98±6      | H            |
| Tamiasius Hudsonicus      | 60.0±15  | 215.3±15  | 0.0073±57 | 4.0±80      | H            |
| Thunnus Albacares         | 80.0±26  | 45349.0±51 | 0.0015±84 | 6.2±47      | H            |
| Tapia glis               | 90.0±27  | 142.0±35 | 0.0024±36 | 3.4±79      | H            |

CFF = critical flicker fusion; Mg = body mass (g); qWg = temperature-corrected (25 °C) mass-specific resting metabolic rate (W/g); light levels: H = high, L = low; NA = no data available for species. Superscript indicates type of measurement: e = electroretinogram; b = behavioural experiments; o = optimum methodology; s = suboptimum methodology; numbers refer to data sources: (1) Crevier & Meister (1998); (2) Adrian & Matthews (1926); (3) Fleishman et al. (1995); (4) Bornshein & McNab (1986); (64) Hudson et al. (1972); (65) Lowe (2001); (66) Jones & Wang (2004). For respiration dynamics of rods and cones (Rubene et al. 2010), we included light levels within our analyses as a categorical variable based on the light conditions experienced by the species during normal activity (i.e. foraging). Species were categorized as inhabiting either high or low light conditions with diurnal terrestrial and nontubid aquatic species coded as inhabiting high light level environments and nocturnal species coded as inhabiting low light levels. As the light levels of species that inhabit turbid waters are typically orders of magnitude lower than typical daylight levels (40–1000 lx; Ali & Klyne 1985; Palmer & Grant 2010; Kreysing et al. 2012) and the barb seal, Pagophilus groenlandicus, regularly forages at depths greater than 200 m (Folkow et al. 2004) where light levels are comparable to nocturnal light levels (Palmer & Grant 2010), we categorized these species as inhabiting low light level environments.

To correct for the phylogenetic nonindependence of species we constructed a composite tree of the study species using published molecular phylogenies and divergence times from various sources (Schoch 1985; Janosy 1986; Mercer & Roth 2003; Hedges et al. 1997; 67) Pauls (1981); 68) Dewar & Graham (1994); 69) Garamszegi et al. (2002); 70) Ivansk & Nelson (2002); 71) Cole & Quiring (1940); 72) Herculano-Houzel et al. (2006); 73) Davenport et al. (2009); 74) Burton (2008); 75) Platel (1979); 76) Aiello & Wheeler (1995); 77) Froese & Pauly (2012); 78) Wallace et al. (2010); 79) Navaret et al. (2011); 80) Meier (1983).

* Indicates species with qWg estimated from swimming speeds extrapolated to zero (see Methods).
2006; Wiens et al. 2006; Benton & Donoghue 2007; Murphy et al. 2007; Brown et al. 2008; Li et al. 2008; Naro-Maciel et al. 2008; Albert et al. 2009; Lim et al. 2010; Little et al. 2010; Perelman et al. 2011; see the Appendix and Fig. A1). In instances in which a divergence time was not available for two species we used the conservatively estimated date of first appearance as the divergence time taken from the Paleobiology Database (Alroy et al. 2008).

As ectotherm metabolic rates vary with temperature, we performed a sensitivity analysis to test the effect of the temperature to which qWg was corrected to by rerunning the main analysis with qWg corrected to both 5 °C and 35 °C (see Appendix). We also carried out a supplemental analysis on a more restricted data set for species with available brain mass data to test for any possible effects of sensory tissue on maximum CFF values (see Appendix).

In total we collected data on maximum CFF, body mass, qWg and light environments for 34 species across the vertebrate classes Elasmobranchii, Actinopterygii, Aves, Amphibia, Reptilia and Mammalia, with further data on brain mass for 28 of these species (Table 1).

**Statistical Analyses**

To test our hypothesis we used a phylogenetic generalized least-squared approach (PGLS) using the caper package (Orme et al. 2012) in R version 2.14.2 (R Development Core Team 2012). The PGLS approach is based on standard generalized least-squared models while also accounting for the nonindependence in the data caused by species’ phylogenetic relationships by incorporating it through the error term structure (Pagel 1999; Rohlf 2001). This error term consists of a matrix of expected trait covariance calculated using the maximum likelihood estimate of lambda (λ), a multiplier of the off-diagonal elements of a phylogenetic variance–covariance matrix that best fits the data. When the data are structured according to a Brownian motion of trait evolution, λ = 1, whereas when the data have no phylogenetic dependency, then λ = 0 (Pagel 1999).

We ran PGLS models with maximum CFF as the response variable, and all combinations of the following explanatory variables: body mass, qWg, light level (high, low) and experimental procedure (ERG, behavioural) with brain mass and methodological optimality included in the sensitivity analysis (see Appendix). We did not include interactions, as there was no a priori reason to include them. We used the Akaike information criterion (AIC), which penalizes extra effective parameters to avoid over-parameterized models, to select the minimum adequate model (Burnham & Anderson 2002).

**RESULTS**

The most parsimonious model (based on AIC) explaining variation in maximum CFF among vertebrates included the terms body mass, log10 of temperature-corrected mass-specific resting metabolic rate (qWg) and light level (Table 2, Tables A1 and A5 in the Appendix). The second most parsimonious model, which fell within two AIC values of the most parsimonious model, retained all tested variables (Table 2). Body mass had a negative effect on the temporal resolution of the sensory system (Table 2, Fig. 2a, Fig. A2 in the Appendix), with a change in body mass of approximately 10 kg resulting in a reduction in CFF of 2 Hz. The metabolic rate of organisms, after correcting for mass, was positively associated with CFF while low environmental light levels were associated with an overall reduction in CFF (Table 2, Fig. 2b, Fig. A2 in the Appendix). Phylogeny was found to have a minimal effect on the resulting models (λ = 0, Table 2) and experimental type was not correlated with CFF (Table 2). Thus, according to our model, small animals with high mass-specific metabolic rates in high light environments possessed the highest maximum CFF and hence greatest ability to perceive temporally dynamic visual information. Conversely, large animals with low mass-specific metabolic rates in low light environments had the lowest CFF.

These results were robust to our sensitivity analysis on both the temperature used to correct ectotherms qWg (taken as 25 °C in the

**Table 2. Coefficients of the two most parsimonious models in the main analysis (based on AIC)**

| Variable | Estimate | SE | t | P |
|----------|----------|----|---|---|
| Model 1 | R²=0.79 |    |   |   |
| Intercept | 118.60 | 11.30 | 10.54 | <0.0001 |
| Mg | $-2 \times 10^4$ | $4 \times 10^3$ | -4.45 | <0.001 |
| log10(qWg) | 13.20 | 4.02 | 3.30 | <0.005 |
| Light.l (low) | -41.12 | 4.87 | -8.44 | <0.0001 |
| Lambda (λ) | Mode | Lower 95% CI | Upper 95% CI |
| Intercept | 0 | 0 | 0.22 |
| Model 2 | R²=0.78 |    |   |   |
| Intercept | 118.90 | 12.00 | 9.94 | <0.0001 |
| Mg | $-2 \times 10^4$ | $4 \times 10^3$ | -4.45 | <0.001 |
| log10(qWg) | 13.24 | 4.08 | 3.24 | <0.005 |
| Light.l (low) | -41.10 | 4.96 | -8.28 | <0.0001 |
| Exp.t (ERG) | -0.51 | 5.08 | -0.10 | 0.92 |
| Lambda (λ) | Mode | Lower 95% CI | Upper 95% CI |
| Intercept | 0 | 0 | 0.22 |

Mg – body mass (g); qWg – temperature-corrected (25 °C in main analysis) mass-specific resting metabolic rate W/g; light.l (low) – effect of low light levels on CFF in comparison to high light levels and exp.t (ERG) – effect of experimental type (ERG – electroretinogram) in comparison to behaviour-based CFF measures.

Figure 2. The effect of (a) body mass (presented on log10 scale) and (b) log10 temperature-corrected mass-specific resting metabolic rate (qWg) on critical flicker fusion frequency (CFF) while controlled for light levels. The minimal adequate model (Results) indicates CFF increases with log10 qWg (13.24 ± 4.08) but decreases with body mass ($-0.0002 \pm 0.00004$). Low light levels are associated with low CFF values ($-41.10 \pm 4.96$) in comparison to high light levels. Figure adjusted to display the intercept at the median value of the unrepresented axis.
main models above; see Methods) and the optimality of study methodology for measuring maximum CFF, with the best models in both sensitivity analyses (based on AIC) including the same terms and trends as found in the main analysis (Tables A2, A3, A5, A6, A7 and A9 in the Appendix). We also found that including brain mass in a restricted data set of 28 species for which brain mass was available did not change the effect of the explanatory variables light levels, qWg and body mass on maximum CFF (Tables A4 and A8 in the Appendix).

**DISCUSSION**

Many of the interspecific and intraspecific interactions that shape species' behaviour and ecology rely on the ability of organisms to process high temporal resolution sensory information. Our results show that, while there is considerable variability in the ability to resolve temporally dynamic visual information across vertebrates, body mass and metabolic rate act as important general constraints on this ability. This is the first study to indicate a general trend in the ability of vertebrates to resolve temporal information; previous studies have generally focused on specific cases of sensory adaptations (Fritsches et al. 2005) and particular environments (Frank 1999; Frank et al. 2012), hence focusing on the particular ecological context of each adaptation or environment. Our findings illustrate the relationship between both physiology and the effects of body mass on the ability to resolve temporal features of the environment on fine timescales, hence linking sensory adaptations to fundamental constraints and trade-offs imposed on all organisms.

Austrom's (1958) hypothesis, that the response dynamics of the retina should be shaped by the organism's particular ecology, predicts that organisms that demand fast visual systems will acquire adaptations increasing CFF values, and hence temporal resolution. For instance, given the strong effect of metabolic rate on CFF, one obvious adaptation is to alter the physiology and metabolism associated with the visual processing systems as seen in the localized heating of tissues in the heads of blowflies (Tatler et al. 2000) and the eyes of predatory swordfish (Fritsches et al. 2005). These tissues increase the temperature around the sensory tissues associated with the blowfly's or swordfish's visual system, which allows for an upregulation of CFF. Similar adaptations are also seen across species of large, fast-swimming predatory billfish (Carey 1982) and Lamnidae sharks (Block & Carey 1985). Physiological adaptations for high-resolution motion detection are also found within specific areas of the retina in some flies, commonly referred to as the 'love spot', which allow them to identify female flight patterns accurately and thus detect mates (Land & Collett 1974). Alterations to the rate of neuron firing, a fundamental limit to the rate of information transfer, through the provision of energy (Laughlin 2001) or changes in the physiological environment, as described above, would also allow for selection on temporal resolution abilities on a neurological level.

In a broader context, it might be expected that manoeuvrability, a vital component of an individual’s ability to respond to the environment, may be one of the main factors determining whether it is necessary to invest in costly temporal information processing. Manoeuvrability, as defined by the ability to change body position or orientation, generally scales negatively with body mass. This negative scaling emerges primarily through the increased inertia and decreased limb stroke rate associated with large body size in both aquatic and volant species (Dudley 2002; Sato et al. 2007; Vogel 2008; Hedrick 2011; Watanabe et al. 2012), while in terrestrial species changes in gait posture that redistribute weight across the limbs can explain such reduced manoeuvrability with body mass (Heglund & Taylor 1988; Biewener 2003). These arguments show that, owing to the laws of physics, larger animals physically respond less quickly to a stimulus. Hence we expect selection against costly investment in sensory systems with unnecessarily high temporal resolution in large animals, as information on such timescales can no longer be utilized effectively. This may explain why larger vertebrates, along with those with low metabolic rates, had lower temporal resolution in our study. This idea is also supported by research showing that faster and more manoeuvrable fly species have higher temporal resolutions (Laughlin & Weckström 1993) and that less manoeuvrable scavenger crabs display slower response dynamics than deeper living predatory species which are likely to have more active lifestyles (Frank et al. 2012).

The effects of body size and metabolic rate on temporal resolution and the presence of sensory adaptations, as discussed above, also point towards an interesting dimension of niche space. Disparity in size and metabolic rate among species within an ecological setting may select for particular sets of adaptations creating a diverse set of sensory systems and interactions. In such a system, species might occupy the same spatial and temporal niche, but could be separated owing to differential responsiveness to environmental signals and cues as a result of having evolved divergent signalling systems along a dimension represented by temporal resolution. For example, it seems at least theoretically possible to encode information in high-frequency signals that can be detected by intended receivers such as conspecifics but that are not susceptible to ‘eavesdropping’ by (generally larger) predators. Ecological systems in which this may be apparent include deep-sea systems where visual signalling is an important determinant of the ability of organisms to interact, and where bioluminescence flashing over wide frequency ranges is ubiquitous (Haddock et al. 2005; Widder 2010).

In conclusion, our results show that the evolution of sensory systems, which play a vital role in ecological interactions, is subject to limitations imposed by metabolic rate and body mass over orders of magnitude in scale. Furthermore, deviations from the expected relationship between temporal perception, body size and metabolic rate are predicted to be subject to selection pressures for physiological, morphological and behavioural adaptations that alleviate these constraints. The generality of these findings suggest that temporal resolution may play a much more important role in sensory ecology than previously indicated, in particular because of its universal effects relating to body size. Further investigations into both the underlying mechanisms of these findings and their importance to ecological functioning are needed.

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APPENDIX

Phylogeny Reconstruction

We used divergence times and phylogenies from the literature to produce a composite phylogeny of the vertebrate species used in our analyses (Fig. A1). For species with no available divergence dates based on molecular data or available published trees, we used conservatively estimated first appearance dates from the Paleobiology Database as an estimate of divergence time (Alroy et al. 2008).

We took divergence dates for the major groups Elasmobranchii, Actinopterygii and Amphibia from the TimeTree database (Hedges et al. 2006). For divergence dates of Carcharhinidae and Sphyridae we used estimates from Lim et al. (2010), while the divergence time between Negaprion brevirostris and Carcharhinus acronotus was estimated based on the appearance of N. brevirostris, the younger of the two species (Negaprion spp.: 30.3 million years ago; Carcharhinus: 46.2 million years ago). For Actinopterygii we used Li et al. (2008) to infer their phylogenetic relationships and divergence times, and Little et al. (2010) for Perciformes’ divergence times. We used Benton & Donoghue’s (2007) estimation of divergence time between anopids (Testudines and Aves) and sauropids (Squamata and Rhynchocephalia) while Perelman et al. (2011) was used for the divergence and phylogenetic relationships among Squamata, Rhynchocephalia, Testudines and Aves. For divergence times within the Squamata we used Wiens et al. (2006), while for Testudines we used Naro-Maciel et al. (2008). We used Brown et al. (2008) for the Aves phylogeny with divergence times between Asio flammeus and Bubo virginianus estimated from the first appearance in the fossil record (Janossy 1986). We used Murphy et al. (2007) for divergence dates of Mammalia orders, while for Primates we used Perelman et al. (2011). Rodentia divergence times were taken from Murphy et al. (2007). All references relating to the phylogeny are given in the reference list.

Sensitivity Analyses

We performed a series of sensitivity analyses to test whether the results of our main analysis were affected by (1) the temperature to which ectoderm species’ metabolic rates were corrected, (2) the inclusion of brain mass as a control for information-processing abilities and (3) the quality of the data used in the analysis.

(1) Ectotherm temperature. We used Q10 values, the fold change in metabolic rate over a temperature change of 10 °C, as defined for each of the major groups (i.e. reptilian, amphibian, etc.; see Methods), to correct ectoderm mass-specific metabolism (qWg) over a temperature range of 5 °C–35 °C. We performed this analysis by rerunning the main analysis with qWg corrected to 5 °C and then corrected to 35 °C. The resulting set of models and the terms that they included are given in Tables A2 and A3. In both analyses the model with the lowest AIC includes the same terms as found in the main analysis, i.e. body mass (Mg), temperature-corrected mass-specific resting metabolic rate (qWg) and light levels, with qualitatively the same significant effects (Fig. A2, Tables A6 and A7).

(2) Brain mass. As the amount of sensory tissue available to an organism may aid in its ability to perceive and process information, brain mass values, measured as wet weight (g), were taken from the literature (Table 1, Methods). As data on brain mass were available for only a subset of 28 species, we included the term brain mass along with the terms used in the main analysis (light levels, qWg, experimental design and body mass) in a series of models performed on the restricted data set (Table A4). While we found a similar trend to the first analysis with a positive effect of log10 mass-specific resting metabolic rate (14.05 ± 4.82) and negative effects of low light levels (−43.02 ± 5.6) and body mass (−0.0002 ± 0.00004), brain mass was found to have no significant effect on CFF levels (Table A8).

(3) Optimality of study methodology for measuring maximum CFF. To ensure that data quality did not affect the results of our analysis we coted the values from each study as either methodologically optimum or methodologically suboptimum. Only data from studies that used a sufficient range of flicker and light intensity within the experimental procedure were coded as methodologically optimum (Table 1, Methods).

To test the possible effect relating to this grouping on our results we ran the main analysis with these categories added as a fixed factor. The resulting set of terms included in the top models are given in Table A5. In the best model based on AIC that includes the methodology optimality term, all terms found in the main analysis (light levels, log10 of qWg and body mass) were found to be included. As in the main results Mg and light levels had significantly negative effects, and log10(qWg) had a significantly positive effect and the methodology optimality term had no significant effect on CFF (Table A9).

As methodologically suboptimum data would be expected to give values below the maximum CFF and hence produce negative residuals in our models, we also performed a Mann–Whitney U test between residuals taken from the main model representing both methodology categories (Table 2, Results in main text). We found no significance difference between the residuals representing methodologically optimum and suboptimum data and also visually found an even spread of each type of residual when plotted (Mann–Whitney U test: $U = 185$, $N_1 = 20$, $N_2 = 14$, $P = 0.09$; Figs A3 and A4).
Figure A1. Species and phylogenetic relationship used in comparative analysis. Scale bar represent 50 million years. See Methods for details.
Figure A2. The effect of body mass (presented on log scale), light levels and log temperature-corrected mass-specific resting metabolic rate (qWg) on critical flicker fusion frequency (CFF). The minimal adequate model (Results) indicates CFF increases with log10 qWg (13.24 ± 4.08) but decreases with body mass (−0.0002 ± 0.00004). Low light levels (blue) are associated with low CFF values (−41.10 ± 4.96) in comparison to high light levels (red). Residual values for each species are shown for different light levels with stems connecting them to the model surface.

Figure A3. Residuals for optimum and suboptimum data quality taken from model 1 in main analysis. Box plot shows median (line), quartiles (box limits), 5th and 95th percentiles (error bars) and outliers (open circles).

Figure A4. Plot of Fig. 1 with data quality represented with methodologically optimum (O) and methodologically suboptimum (S) data. Slopes corrected to represent the intercepts of each explanatory variable at the median value of (a) log10(qWg) and (b) log10(Mg).

Table A1

| Model | Explanatory variables in model | AIC | AICΔ |
|-------|--------------------------------|-----|------|
| 1     | Mg, log10(qWg)                  | 275.70 | 0 |
| 2     | Mg, log10(qWg)                  | 277.68 | 1.98 |
| 3     | Mg, log10(qWg)                  | 283.94 | 8.24 |
| 4     | Mg, log10(qWg)                  | 285.91 | 10.21 |
| 5     | Mg, log10(qWg)                  | 291.56 | 15.86 |
| 6     | Mg, log10(qWg)                  | 293.41 | 17.71 |
| 7     | Mg, log10(qWg)                  | 298.90 | 23.20 |
| 8     | Mg, log10(qWg)                  | 300.63 | 24.93 |
| 9     | Mg, log10(qWg)                  | 315.06 | 39.36 |
| 10    | Mg, log10(qWg)                  | 315.07 | 39.37 |
| 11    | Mg, log10(qWg)                  | 316.84 | 41.14 |
| 12    | Mg, log10(qWg)                  | 316.92 | 41.22 |
| 13    | Mg, log10(qWg)                  | 320.67 | 44.97 |
| 14    | Mg, log10(qWg)                  | 322.67 | 46.97 |
| 15    | Mg, log10(qWg)                  | 324.45 | 48.75 |

Mg – body mass (g); qWg – temperature-corrected (25 °C) mass-specific resting metabolic rate W/g; AIC – Akaike’s information criterion. AICΔ gives the difference between each model AIC and that of the lowest AIC found for any model. Terms retained are represented with + symbols, while terms not retained are represented by – symbols.

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Table A2
Terms included in models performed in analysis with mass-specific resting metabolic rate qWg corrected to 5 °C

| Model | Explanatory variables in model | AIC  | AICΔ |
|-------|-------------------------------|------|------|
| 1     | + + + -                        | 274.67 | 0 |
| 2     | + + + +                        | 276.61 | 1.94 |
| 3     | + + - +                        | 283.94 | 9.27 |
| 4     | + - - +                        | 285.91 | 11.24 |
| 5     | - - + +                        | 289.58 | 14.91 |
| 6     | - - - +                        | 291.58 | 16.91 |
| 7     | - - - -                        | 298.90 | 24.23 |
| 8     | + - - -                        | 300.63 | 25.96 |
| 9     | + + + -                        | 314.60 | 39.93 |
| 10    | + + - +                        | 315.01 | 40.34 |
| 11    | + - - -                        | 316.58 | 41.91 |
| 12    | + + + -                        | 317.01 | 42.34 |
| 13    | + + - +                        | 320.67 | 46.00 |
| 14    | + - - +                        | 322.67 | 48.56 |
| 15    | - - - -                        | 324.45 | 49.78 |

Mg = body mass (g); AIC = Akaike’s information criterion. AICΔ gives the difference between each model AIC and that of the lowest AIC found for any model. Terms retained are represented with + symbols, while terms not retained are represented by symbols.

Table A3
Terms included in models performed in analysis with mass-specific resting metabolic rate qWg corrected to 35 °C

| Model | Explanatory variables in model | AIC  | AICΔ |
|-------|-------------------------------|------|------|
| 1     | + + + -                        | 277.39 | 0 |
| 2     | + + + +                        | 279.26 | 1.87 |
| 3     | + + - +                        | 283.94 | 6.55 |
| 4     | + - - +                        | 285.91 | 8.52 |
| 5     | - - + +                        | 294.26 | 16.87 |
| 6     | - - - +                        | 295.79 | 18.40 |
| 7     | - - - -                        | 298.90 | 21.51 |
| 8     | + - - +                        | 300.63 | 23.24 |
| 9     | + + + -                        | 315.97 | 38.58 |
| 10    | + + - +                        | 316.49 | 39.10 |
| 11    | + - - - +                      | 317.53 | 40.14 |
| 12    | + - - - -                      | 317.89 | 40.50 |
| 13    | + - - - +                      | 320.67 | 43.28 |
| 14    | + - - - -                      | 322.67 | 45.28 |
| 15    | - - - - -                      | 324.45 | 47.06 |

Mg = body mass (g); AIC = Akaike’s information criterion. AICΔ gives the difference between each model AIC and that of the lowest AIC found for any model. Terms retained are represented with + symbols, while terms not retained are represented by symbols.

Table A4
Terms included in models performed in analysis including brain mass as a factor

| Model | Explanatory variables in model | AIC  | AICΔ |
|-------|-------------------------------|------|------|
| 1     | + + + -                        | 219.56 | 0 |
| 2     | + + + +                        | 221.27 | 1.71 |
| 3     | + + - +                        | 221.54 | 1.98 |
| 4     | + - - +                        | 223.26 | 3.70 |
| 5     | - - + +                        | 226.27 | 6.71 |
| 6     | - - - +                        | 227.99 | 8.43 |
| 7     | + - - -                        | 228.24 | 8.68 |
| 8     | + + + -                        | 229.94 | 10.38 |
| 9     | + + - +                        | 234.55 | 14.99 |
| 10    | + - - +                        | 235.91 | 16.35 |

Mg = body mass (g); qWg = temperature-corrected (25 °C) mass-specific resting metabolic rate W/g; AIC = Akaike’s information criterion. AICΔ gives the difference between each model AIC and that of the lowest AIC found for any model. Terms retained are represented with + symbols, while terms not retained are represented by symbols.

Table A5
Terms included in models performed in analysis with mass-specific resting metabolic rate qWg corrected to 35 °C

| Model | Explanatory variables in model | AIC  | Methodology |
|-------|-------------------------------|------|-------------|
| 1     | + + + +                        | 275.05 | 0            |
| 2     | + + + +                        | 275.80 | 0.75         |
| 3     | + + + +                        | 283.74 | 8.69         |
| 4     | + + + +                        | 283.90 | 8.85         |
| 5     | + + + +                        | 291.74 | 16.69        |
| 6     | + + + +                        | 293.69 | 18.64        |
| 7     | + + + +                        | 298.86 | 23.81        |
| 8     | + + + +                        | 300.49 | 25.34        |
| 9     | + + + +                        | 315.35 | 40.30        |
| 10    | + + + +                        | 315.37 | 40.32        |
| 11    | + + + +                        | 317.01 | 41.99        |
| 12    | + + + +                        | 317.29 | 42.24        |
| 13    | + + + +                        | 320.92 | 45.87        |
| 14    | + + + +                        | 322.26 | 47.21        |
| 15    | + + + +                        | 323.38 | 48.33        |

Mg = body mass (g); methodology – optimality of study methodology; AIC = Akaike’s information criterion. AICΔ gives the difference between each model AIC and that of the lowest AIC found for any model. Terms retained are represented with + symbols, while terms not retained are represented by symbols.

Table A6
Coefficients of the best 5 °C model (based on AIC) for addition analysis with qWg corrected to 5 °C

| Variable | Estimate | SE  | t    | P    |
|----------|----------|-----|------|------|
| 5 °C model |          |     |      |      |
| Intercept | 110.3    | 8.45| 13.05| <0.0001 |
| Mg log₁₀(qWg) | -2 x 10^4 | 4 x 10^5 | -4.40 | <0.001 |
| Light.1 (Low) | 9.5      | 2.73| 3.48 | <0.005 |
| Lambda (λ) | 41.2     | 4.78| -8.62| <0.0001 |
| Mode Lower 95% CI | Upper 95% CI | 0.21 |

Mg = body mass (g); qWg = temperature-corrected (5 °C) mass-specific resting metabolic rate W/g; light.1 = effect of low light levels on CFF in comparison to high light levels.

Table A7
Coefficients of the best 35 °C model (based on AIC) for both the main analysis and each of the sensitivity analyses

| Variable | Estimate | SE  | t    | P    |
|----------|----------|-----|------|------|
| 35 °C model |          |     |      |      |
| Intercept | 122.5    | 13.7| 8.97 | <0.0001 |
| Mg log₁₀(qWg) | -2 x 10^4 | 4 x 10^5 | -4.72 | <0.0001 |
| Light.1 (Low) | 15.23    | 5.11| 2.98 | <0.01 |
| Lambda (λ) | -41.4    | 4.97| -8.30| <0.0001 |
| Mode Lower 95% CI | Upper 95% CI | 0.25 |

Mg = body mass (g); qWg = temperature-corrected (25 °C in main analysis) mass-specific resting metabolic rate W/g; light.1 = effect of low light levels on CFF in comparison to high light levels.

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Table A8

Coefficients of the model including brain mass using reduced data set of \( N = 28 \)

| Variable          | Estimate | SE  | \( t \)  | \( P \)   |
|-------------------|----------|-----|---------|----------|
| Brain model       | 122.0    | 13.22 | 9.23    | <0.0001  |
| \( R^2 = 0.78 \)  |          |      |         |          |
| Intercept         | 2 \times 10^4 | 4 \times 10^3 | -4.33   | <0.001   |
| Mg                | 14.05    | 4.82 | 2.91    | <0.001   |
| \( \log_{10}(qWg) \) | -43.02  | 5.60 | -7.69   | <0.0001  |
| Light.l (Low)     | -0.005   | 0.01 | -0.49   | 0.63     |
| Brain mass        |          |      |         |          |
| Mode              | 0        | 0    | 0       | 0.30     |
| Lower 95% CI      |          |      |         |          |
| Upper 95% CI      |          |      |         |          |

Mg – body mass (g); \( qWg \) – temperature-corrected (25°C in main analysis) mass-specific resting metabolic rate W/g; light.l – effect of low light levels on CFF in comparison to high light levels; brain mass (g).

Table A9

Coefficients of the model including optimality of methodology factor

| Variable                               | Estimate | SE  | \( t \)  | \( P \)   |
|----------------------------------------|----------|-----|---------|----------|
| Model including optimality of study methodology \( R^2 = 0.79 \) |          |      |         |          |
| Intercept                              | 124.59   | 11.6 | 10.7    | <0.0001  |
| Mg                                     | -2 \times 10^4 | 4 \times 10^3 | -4.92   | <0.001   |
| \( \log_{10}(qWg) \)                   | 13.8     | 3.94 | 3.50    | <0.005   |
| Light.l (Low)                          | -43.1    | 4.89 | -8.81   | <0.0001  |
| Method (optimal)                       | -7.68    | 4.917| -1.56   | 0.13     |
| Lambda (\( \lambda \))                | 0        | 0    | 0       | 0.22     |

Mg – body mass (g); \( qWg \) – temperature-corrected (25°C in main analysis) mass-specific resting metabolic rate W/g; light.l – effect of low light levels on CFF; method – optimality of study methodology.