#### 1 How has the environment shaped geographical patterns of insect body sizes? A test 2 of hypotheses using sphingid moths 3 Nicolas Beerli<sup>1</sup>, Florian Bärtschi<sup>1,2</sup>, Liliana Ballesteros-Mejia<sup>3</sup>, Ian. J. Kitching<sup>4</sup> & Jan 4 Beck<sup>1,5\*</sup> 5 6 7 1) University of Basel, Department of Environmental Science, Basel, Switzerland 8 2) Umweltbüro Bärtschi, Basel, Switzerland 9 3) Muséum national d'Histoire naturelle, Sorbonne Université, Institut de Systématique, Évolution, 10 Biodiversité (ISYEB). UMR 7205 - CNRS, MNHN, UPCM, EPHE, Paris, France 11 4) Department of Life Sciences, Natural History Museum London, London, UK 12 5) University of Colorado, Museum of Natural History, Boulder, USA 13 \*) Correspondence: Jan Beck, University of Colorado, Museum of Natural History, 265 UCB, 14 Boulder, CO 80309, USA. E-mail: jan.beck@colorado.edu 15 16 Manuscript for Journal of Biogeography, Humboldt special issue 17 18 Research paper 19 20 Short title: Geographical patterns of moth body sizes 21 **Keywords:** Hawkmoths; Lepidoptera; Bergmann's rule; comparative; ectotherms; 22 phylogeny; 23 Word count: Abstract 300 words; main text 5404; 4 figures, no tables; 56 references; 24 25 Acknowledgements 26 We thank Walter Jetz for fruitful discussions on the subject, and Meret Hornstein, Katherine 27 Winkel, Jonas Egli and Gunnar Brehm for constructive comments on earlier drafts. Blanca Huertas 28 supported our measuring of museum specimens. We received a travel grant from the Synthesys 29 program of the European Community Research Infrastructure Action (FP7 "Capacities" program). 30 This study was funded in part by NERC grant number NE/P003915/1 to IJK, the modelling of 31 species ranges by Swiss National Science Foundation grant 3100AO 119879 to JB. 32 33 **Biosketch**

- 34 Nicolas Beerli completed his MSc in the research team of JB, which focusses on insect
- 35 macroecology and the environmental impacts on biodiversity. *Author contributions*: NB and JB
- designed the study and analysed data, supported by FB; JB, NB, LB-M and IJK provided sphingid
- 37 moth data; NB, JB and IJK wrote the manuscript, with input from all authors.
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# 40 ABSTRACT

- 41 Aim: We mapped the geographical pattern of body sizes in sphingid moths and
- 42 investigated latitudinal clines. We tested hypotheses concerning their possible
- 43 environmental control, i.e., effects of temperature (negative: temperature size rule or
- 44 Bergmann's rule; positive: converse Bergmann rule), food availability, agility in densely
- 45 vegetated habitats, robustness to starvation during extreme weather, and seasonality.
- 46 Location: Old World and Australia/Pacific region
- 47 **Methods:** Body size data of 950 sphingid species were compiled and related to their
- 48 distribution maps. Focusing on body length, we mapped the median and maximum size
- 49 of all species occurring in 100 km grid cells. In a comparative approach we tested the
- 50 predictions from explanatory hypotheses by correlating species' size to the average
- 51 environmental conditions encountered throughout their range, under univariate and
- 52 multivariate models. We accounted for phylogeny by stepwise inclusion of
- phylogenetically informed taxonomic classifications into hierarchical random-intercept
   mixed models.
- 55 **Results:** Median body sizes showed a distinctive geographical pattern, with large species
- 56 in the Middle East and the Asian tropics, and smaller species in temperate regions and the
- 57 Afrotropics. Absolute latitude explained very little body size variation, but there was a
- 58 latitudinal cline of maximum size. Species' median size was correlated to net primary
- 59 productivity, supporting the food availability hypothesis, whereas support for other
- 60 hypotheses was weak. Environmental correlations contributed much less (i.e., <10%) to
- 61 explaining overall size variation than phylogeny (inclusion of which led to models
- 62 explaining >70% of variability).
- 63 **Main conclusion:** The intuitive impression of larger species in the tropics is shaped by
- 64 larger size maxima. Median body sizes are only very weakly related to latitude. Most of
- 65 the geographic variation in body size in sphingid moths is explained by their
- 66 phylogenetic past. NPP and forest cover correlate positively with the body size, which
- 67 supports the idea that food availability allowed the evolution of larger sizes.
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- 69

#### 70 INTRODUCTION

71 Tropical insects of impressively large body size compared to their temperate 72 counterparts are a common observation of field naturalists, as well as those admiring 73 museum collections - starting with A. v. Humboldt's scientific explorations of the Neotropics around 1800, on which he collected insects of appreciable size (Barragán et 74 75 al., 2009). 50 years later his successor, Amazonian explorer H.W. Bates (1864, p.115) reported accidentally shooting hawkmoths so large he mistook them for small birds. 76 77 However, it remains unclear whether such observations necessarily imply a larger size for 78 the average tropical insect species – it just may be that their maxima are more extreme in 79 regions of higher species richness, and it is these that capture our attention (cf. coloration 80 in tropical birds; Bailey, 1978). Furthermore, it must be expected that body sizes, like 81 species richness or other trait variation, exhibit more complex geographical patterns than simply a latitudinal gradient (Hawkins & Diniz-Filho, 2004). Here, using sphingid moths 82 83 (hawkmoths) as the exemplar, we map for the first time the body size distribution of a 84 group of insects at intercontinental geographic scale (i.e., global excluding the Americas) and at high spatial resolution. In a comparative approach we investigate how 85 environmental conditions and phylogenetic inertia may have shaped these patterns. 86

87 Body size is a prime example of geographic trait variation, with important links to 88 physiological functioning, population-level processes and biodiversity as a whole 89 (Gaston & Blackburn, 2000; Blanckenhorn, 2000; Brown et al., 2004). Consequently, its 90 relationships to environmental variables are of major interest (Gaston & Chown, 2013), 91 yet conclusive, large-scale studies, intra- or interspecific, have mostly been carried out on 92 endotherms (Blackburn & Hawkins, 2004; Freckleton et al., 2003; Olson et al., 2009). 93 Environmental gradient studies on ectotherm body sizes are scarcer, less firmly rooted in 94 theory, and provide more conflicting results at an intraspecific (Ashton et al., 2003; Puzin 95 et al., 2014) as well as an interspecific level (Entling et al., 2010; Fattorini et al., 2013; 96 Slavenko & Meiri, 2015; Zeuss et al., 2017; Brehm et al., 2019; Pallares et al., 2019).

97 The formulation of eco-geographical rules has been the cornerstone of attempts to 98 describe patters and understand mechanisms of trait variation, and the most well-known 99 of these, Bergmanns' rule (Meiri, 2011) concerns body size. Originally primarily addressing endothermic vertebrates, it proposes an interspecific body size increase 100 towards higher latitudes if restricted to closely related taxa. While the pattern itself is 101 102 well-supported (Meiri & Dayan, 2003), fewer studies have tested and supported the 103 underlying thermoregulatory mechanism (Blanckenhorn & Demont, 2004; Watt et al., 104 2010; Meiri, 2011; Fattorini et al., 2013; Beck et al., 2016). Other mechanistic, 105 environmental effects on body size, both evolutionary and developmental, have been suggested for insects and other ectotherms (Chown & Gaston, 2010, 2013; Chown & 106 107 Nicolson, 2013), and these may not act in a mutually exclusive manner. Furthermore, body sizes are strongly affected by phylogeny, which complicates analyses and the 108 109 inference on environmental causes of the observed variation (Freckleton & Jetz, 2009).

110 Using data on size, high-resolution maps of species' geographical ranges, and a 111 phylogenetically-informed taxonomic classification, we mapped the geographical body 112 size pattern of sphingid moths and tested its consistency with the following hypotheses 113 while accounting for phylogenetic effects.

(1) Bergmann's rule expects a negative relationship between body size and
environmental temperature. It assumes that thermoregulation, through the ratio of
heat-exchanging surface and heat-producing volume, favors larger animals in colder
climates. Although it was originally focused on endotherms, Zamora-Camacho *et al.*(2014) argued for a Bergmann-type heat preservation mechanism in an ectotherm
lizard. Some studies have also investigated Bergmann's rule in Lepidoptera, with

- 120 mixed results (e.g., Beck *et al.*, 2016, Brehm *et al.*, 2019). Furthermore, adult
- sphingid moths are known for pre-flight thoracic muscle temperature regulation
- 122 (Heinrich 1993), hence temperature efficiency may be relevant. Alternatively, an
- 123 intraspecific effect known as the 'temperature size rule' (TSR; Kingsolver & Huey,
- 124 2008; Chown & Gaston, 2010) predicts the same pattern. Laboratory experiments
- 125 with many ectotherms have shown that individuals develop slower yet mature into a
- 126 larger body size under colder temperatures (Atkinson, 1994). This may be either an
- 127 effect of adaptive plasticity or it may be due to non-adaptive constraints of
- temperature on growth rates (e.g., the discrepancy between oxygen supply and
- demands; see Makarieva *et al.*, 2005 for an interspecific approach). The TSR has also
  been shown to apply to a sphingid moth (*Manduca sexta*; Davidowitz & Nijhout,
- 131

2004).

- (2) The 'converse Bergmann pattern' expects a positive relationship of body size and
  temperature. The reasoning here is that lower temperatures lead to lower growth rates
  in ectotherms, favoring species that become adult (i.e., reproductive) at smaller size
  (Meiri, 2011; Shelomi, 2012).
- (3) The resource availability hypothesis postulates an increase of body size with
  increased food availability (Blackburn *et al.*, 1999; Watt *et al.*, 2010).
- 138 Intraspecifically, lower food availability is known to result in smaller specimens
- (Slansky & Scriber, 1985). If this mechanism was relevant interspecifically, we can
  predict a body size increase with net primary productivity, which we use as a proxy
  for food availability in our herbivorous study taxon.
- (4) The starvation hypothesis postulates that larger species occur in regions with a higher
  risk of unforeseeable starvation. With increasing body size fat storage increases faster
  than metabolic rate, hence survival of such catastrophic events becomes more likely
  (Chown & Gaston, 2010). This hypothesis assumes that insect species in variable
  climates need more physiological tolerances than those in zones of stable climate
  (Addo-Bediako *et al.*, 2000). We predict that body size increases with increasing
  temperature extremes during their active period.
- (5) The seasonality hypothesis (Mousseau, 1997) assumes that in seasonal habitats,
  where a part of the year is unsuitable for growth, smaller species profit from a faster
  completion of their reproductive cycle. We expect a negative link of seasonality and
  body size.

Phylogeny plays an important role in the current spatial distribution of species' traits, due to prior adaptation to past environmental conditions and the constraints of evolutionary history (Gaston & Chown, 2013). It can be assumed that a part of the variation in body size is explained by phylogenetic inertia. For interspecific comparisons, such phylogenetic signals within the data must be accounted for to avoid spurious effects on the analysis of environmental predictor variables (Freckleton & Jetz, 2009, and references therein).

160 Here, we first map geographical patterns of size in sphingids moths at large 161 geographical scale by combining detailed distribution data and body size measurements. 162 We explored whether hawkmoths are larger or smaller towards the equator using a simple latitudinal cline. We then tested the above hypotheses by investigating the link between 163 164 body size and environmental variables, such as temperature, seasonality, forest cover, 165 temperature extremes and net primary productivity (NPP). We used a comparative, 166 'species-focused' approach where every species counts equally (Chown & Gaston, 2010), 167 which is evolutionarily more informative than a 'geography-focused' approach (i.e., 168 comparing grid cells). The latter would be overly impacted by widespread species that 169 occur in many cells (Jetz & Rahbek, 2002). Phylogenetic information was included into

- 170 the analyses at different taxonomic classification levels, which helped to identify the
- 171 importance of phylogenetic history in comparison to environmental effects.
- 172

#### 173 METHODS

174 Hawkmoths or sphingids are a family of Lepidoptera that have attracted the 175 attention of insect collectors for centuries (Kitching & Cadiou, 2000). Consequently, they 176 are more extensively studied, taxonomically and biogeographically, than most other 177 insects. Recently, Ballesteros-Mejia et al. (2017) published and analyzed detailed 178 geographic ranges for all species occurring outside the Americas (data available at 179 www.mol.org). These maps stem from expert-edited species-distribution modelling 180 (relating published and unpublished occurrence records to climate and vegetation 181 variables, at 5 x 5 km resolution). Details on the procedures of modelling and validation of geographic range maps are found in Ballesteros-Mejia et al. (2017). Of 981 species 182 183 occurring in the research region, we considered 947 species for the present analysis. The 184 reasons for excluding some taxa were phylogenetic uncertainty (i.e., unreliable species 185 status), missing environmental data (small-island endemics of the far eastern Pacific) or 186 because no male specimens (or images) were available for body size measurement (see 187 below for a further reduction to 938 species for body length measures).

188 Theoretically, body mass is the physiologically most meaningful size metric 189 (Gaston & Blackburn, 2000), but it is difficult to measure when relying on dried museum 190 specimens or images. Many length measurements have been shown to correlate with 191 body mass and such linear measurements are commonly used as a proxy for body size in 192 insects (Chown & Gaston, 2010; Brehm et al., 2019). We compiled data on body length 193 (head to tip of abdomen), thorax width, and forewing length (base to tip). Measurements 194 were taken from scaled pictures published in d'Abrera ([1987]; 459 species), from images 195 that are publicly available (28 species), and unpublished colour photographs (38 species, 196 IJK's personal photographic print collection). The remaining 422 species, which were not 197 available in scaled illustrations, were measured from pinned specimens in the collection 198 of the Natural History Museum, London. Details of the measured specimens can be found 199 in Appendix ES1. All measurements were taken with a digital caliper, rounded to 0.1 200 mm. Since there is sexual size dimorphism in some sphingid groups (e.g., tribes 201 Smerinthini, Ambulycini, where males tend to be smaller than females; Kitching & 202 Cadiou, 2000) we measured and compared only males (the more common sex in 203 collections) for consistency. In other Lepidoptera taxa, size measures for males and 204 females are highly correlated (data from Brehm et al., 2019, see there also for further 205 discussion). Furthermore, we would note that any resulting size underestimates within 206 those clades' size would be controlled for by fitted random effects models for 207 phylogenetic association (in mixed models, see below).

As in other macro-studies (e.g., Zeuss *et al.*, 2017) we were only able to measure 208 209 one specimen per species (often only one picture or specimen was available). Although 210 body size can be variable within species, such a sample (n = 1) is an unbiased estimate of 211 the mean. Although a single-specimen measure would be unsuitable for intraspecific 212 studies, it should vield near-identical patterns to averages based on many specimens if 213 studied across many species. This is because randomly-distributed errors will even out. 214 and correlation analyses will provide reliable results. We confirmed this theoretical 215 expectation with subsampling simulations based on moth body size data from Brehm et 216 al. (2019; J. Beck & G. Brehm, unpubl.). Furthermore, we assessed the size variability in eight abundant sphingid species, where the body lengths of 208 specimens (14-34 per 217 218 species) were measured from scans of collection drawers (Johnson et al., 2013; Trueman 219 & Yeates, 2015; data in ES1). The absolute difference between mean body lengths of

these specimens and the (independent) single specimen-measure was on average 3.4 mm,

- which we find small when considering an interspecific body length range of 55.8 mm
- 222 (from 12.5 to 68.3 mm) in our dataset. Thus, while intraspecific effects may not be
- 223 entirely negligible, they are probably irrelevant in comparison to interspecific variability

in a dataset with many hundreds of species. Over our large sample of 938 species they

- 225 may just add some additional random noise. Furthermore, intraspecific variability in
- Lepidoptera often appears large due to occasional 'dwarf specimens', but these do not
- tend to be illustrated in pictures whenever 'normal' specimens are available. Measuring tens of thousands of specimens to fully appreciate species' intraspecific variation was
- simply beyond of the scope of this research project. Data for the size measures for all
- 230 species in analysis are available in ES1.
- 231
- 232 Environmental predictors

To test our hypotheses on the environmental control of body size, we considered the mean annual temperature of frost-free months, net primary production, forest cover, extreme temperatures and seasonality. Additionally, absolute latitude was used as a nonenvironmental predictor.

- Mean annual temperature was calculated for those months with an average monthly
   temperature above 0°C. Data were based on monthly temperature data from the
   WorldClim database (resolution 30 arc-second, time period 1950-2000; Hijmans et
   al., 2005). Excluding cold winter months is reasonable because sub-zero winter
   temperatures are quite irrelevant for diapausing organisms while including them into
   averages would bias relevant temperatures downward.
- Net primary production (NPP) data was taken from remotely sensed normalized
  differential vegetation index values (NDVI), corrected for modelled periods of water
  limitation (Running *et al.*, 2004; time period 2000-2014). In this dataset 'No Data'
  values caused by lack of green reflection were set to zero where they occurred on
  land, as vegetation-free landscapes (e.g., deserts, glaciers) must be interpreted as
  unproductive for our purposes, rather than being excluded from analysis.
- *Forest cover* (in percent) was based on Tuanmu & Jetz (2014), who provided 12
   generalized land cover maps based on a consensus land cover dataset. Four of these
   land cover categories were forests (evergreen/deciduous needleleaf trees; evergreen
   broadleaf trees; deciduous broadleaf trees, mixed/other trees). Percentages from these
   four forest types were summed (and corrected to 100% where this value was
   exceeded due to integer rounding).
- 255 *Temperature extremes* data were based on temperature data from Smith *et al.* (2008) 0 256 at a resolution of 5°. For every grid cell, data included monthly temperature deviation of a long-term average (128 years: 1880-2008). In the northern hemisphere, we 257 excluded data north of 35°N for November, December, January and February to 258 259 account for diapause; analogously we excluded data south of 35°S for May, June, 260 July and August in the southern hemisphere. We then counted within each grid cell the months with temperature values that were below one standard deviation (SD) 261 from the grid cell mean (referring to SD of all grid cells in analysis). The number of 262 263 months was divided by the estimated length of the hawkmoths' activity period (nine months above 35° S/N and 12 in the other areas) to render them comparable. This 264 provided a metric of extreme negative temperature events (i.e., 265

266  $\frac{\text{months}|\text{ temp. } \leq \text{mean} - \text{SD}}{\text{months of activity per year}}$  in 128 years of data). Finally, the data were interpolated to a 267 finer resolution using inverse distance weighting (search radius 6 points) to make

267 finer resolution using inverse distance weighting (search radius 6 points), to make

- data comparable with other environmental data. The higher our metric, the moremonths with extremely low temperatures occurred.
- *Temperature seasonality* data were used from the WorldClim database (Hijmans *et al.*, 2005). It is the standard deviation of the annual temperature seasonality, multiplied by 100.

Predictor data were re-projected to a Mollweide World equal area projection (bilinear resampling) to match moth data. Pixels in oceans, seas or lakes >1 km<sup>2</sup> were excluded from all raster datasets. We carried out the handling and extraction of spatial data in ArcGIS 10.3 (http://desktop.arcgis.com), subsequent data compilation and all further analyses were done in R 3.3 (https://cran.r-project.org/; packages *ade4, ape, caper, data.table, ecodist, lme4, matrix, nlme, xtable*).

279 We used two different approaches to map interspecific body size clines. In an 280 assemblage-based (Chown & Gaston, 2010) (or grid cell-based) approach we mapped the 281 median body length of all species occurring in 100 x 100 km grid cells. Grid-cells containing  $\leq 5$  species were removed to reduce random noise. We measured the 282 283 variability of body length with the interquartile range, as well as the maximum body length (i.e., largest species per cell). We used this approach for map visualizations and 284 for assessing body size variation with (absolute) latitude (using adequate techniques to 285 286 account for spatial autocorrelation in statistical tests, details in ES3). However, our main 287 analyses did not follow this grid cell-based approach as it is weakened by not accounting for phylogenetic effects, and by pseudo-replication due to one species occurring in 288 289 several (or many) grid cells (Jetz & Rahbek, 2002; Entling et al., 2010).

290 In our comparative approach (individual-species focus) we treated each species as 291 one data point. This also allowed inclusion of phylogenetic effects into models. To 292 associate environmental predictors with each species, for all environmental variables 293 (except forest cover) we calculated the mean across those 5 x 5 km pixels where the 294 species occur (according to our range maps). For forest cover, we calculated the median 295 at point localities of raw distribution records of high spatial accuracy (<1 km uncertainty) 296 to minimize error on habitat association, excluding ten species with  $\leq 5$  distribution 297 records.

298

#### 299 Data analyses

300 We replaced 24 'No Data' values for forest cover by the means of the variable so as not to lose these species from multivariate models (Zuur et al., 2010). Furthermore, all 301 data were standardized with a z-transformation  $\left(\frac{x-\mu}{\sigma}\right)$ , where  $\mu$  is the mean and  $\sigma$  is the 302 303 standard deviation) to make model coefficients comparable across different analyses. 304 Body sizes, temperature extremes and seasonality were log<sub>10</sub>-transformed after visual 305 inspection of histograms to reach normality prior to the z-transformations, whereas 306 temperature data was  $x^3$ -transformed. NPP data (already normally distributed) and forest 307 cover data (percent values) were standardized without transformation. We assured the 308 fulfillment of model assumptions by checking visually for residual normality, outliers 309 and heteroscedasticity (using R diagnostic plots).

310 Statistical analyses were conducted for all three body size measures (forewing 311 length, thorax width, body length) as well as the product of body length *x* thorax width as 312 response variables, but we present only body length analyses in the main text. This 313 variable is less affected by different body shapes among subfamilies (e.g., systematic 314 differences in thorax/wing length ratios). Models with body length as a response also 315 received the highest pseudo- $r^2$ s in environmental models. We report results for the other 316 metrics in the Electronic Supplement (ES5). In some specimens, no body lengths could

317 be measured due to missing abdomens (a common issue in Lepidoptera collections where

318 abdomens may be removed for genitalia dissection or simply be lost over time), which

319 reduced sample size for these analyses to 938 species.

320

#### 321 Phylogenetic effects

322 It must be assumed that a substantial part of body size variation is due to 323 phylogenetic inertia – i.e., closely related species are similar in size. Such phylogenetic 324 signal must be accounted for in interspecific comparisons to avoid spurious effects of 325 non-independent data. We employed a stepwise, hierarchical approach to do so, 326 accounting for the fact that we had a reliable, phylogenetically informed taxonomic 327 classification, but not a true phylogeny with branch lengths, available for analyses (this would be required for many other approaches). Our classification was based on the 328 329 Sphingidae Taxonomic Inventory (http://sphingidae.myspecies.info), which builds on the 330 tribal-level molecular phylogeny of Kawahara et al. (2009) and integrates most recent 331 findings of phylogenetic and taxonomic studies, molecular and morphology-based. The 332 classification is available in ES2.

333 We started with environmental models containing no phylogenetic information, 334 using a generalized least squares (gls) model. In three further models, we added 335 information on subfamily, tribe and genus associations stepwise into hierarchical linear 336 mixed effect models (lme) as random intercept effects, whereas links with the 337 environment were modelled as fixed effects (Zuur, 2009; see Stone et al. (2011) for mixed models as a method to account for phylogenetic effects). To assess the effect of 338 339 phylogeny onto body size variability, we compared model Akaike information criteria 340 (AICs) and pseudo- $r^2$  values (i.e., correlation of predicted vs. observed values), as well as the standardized coefficients fitted for environmental effects. For the model without 341 342 random effects, we had to use a different algorithm (gls) to that used for models with 343 random effects (*lme*). However, the *gls* and *lme* model coefficients are comparable 344 (Pinheiro & Bates, 2006).

- 345
- 346 Univariate models

We correlated body length in separate models with every environmental predictor variable, using the stepwise approach described above to account for phylogeny. All univariate *lme* analyses were conducted with restricted maximum likelihood estimation (REML), since this is required for a comparison between models with differently nested random structure (Zuur, 2009).

- 352
- 353 Multivariate models

Multivariate analyses were conducted to investigate the independent influence of each predictor variable, using all predictors except latitude. Unlike for univariate *lme* models, we used maximum likelihood (ML) fitting for multivariate *lme* models because AICs from models with a different structure of fixed effects can only be compared when based on ML fits (even though REML fits are less biased; Zuur, 2009). With large sample sizes, differences in models fitted with the two different methods diminish (Zuur, 2009).

To detect and account for collinearity among the predictor variables, the variance
 inflation factor (VIF) of the multivariate model with all predictor variables was

363 calculated. Zuur *et al.* (2010) recommend dropping the predictor variable with the highest

364 VIF, then recalculating the VIF values with the new model, repeating this until the VIF

365 values are <3.

366

#### 367 **Results**

368 There is no clear latitudinal cline in median body lengths (Fig. 1A). Rather, 369 species in the Arabian Peninsula, in South Asia and in the Indo-Australian tropics are 370 distinctively larger than temperate as well as Afrotropical taxa. A grid cell-based 371 correlation of median BL with distance from the equator yields  $r^2 = 0.19$ , but it is non-372 significant due to high spatial non-independence of data (ES3, Fig. S3.4). However, when 373 comparing temperate Eurasia and Africa it becomes clear that the variability in body 374 lengths is larger in the Afrotropics, i.e., there are smaller and larger species than in 375 temperate regions, leading to similar medians. Variability (Fig. 1B) is also high in desert 376 regions of Africa and Asia, despite relatively low species richness (not shown; 377 Ballesteros et al., 2017). Interquartile ranges are significantly negatively correlated with distance from the equator ( $r^2 = 0.47$ ; ES3, Fig. S3.5), and maximum body length per grid 378 379 cell) is strongly and significantly negatively correlated to distance from the equator ((Fig. 380 1C;  $r^2 = 0.54$ , ES3, Fig. S3.6). A map of minimum BL (ES3, Fig. S3.3) shows that the 381 tropics also have smaller species than temperate regions, which supports the assessment 382 that the latitudinal variation of maximum body size is mainly a function of greater 383 variability in more species-rich regions (i.e., larger sample sizes).

Plotting median data in a comparative approach (Fig. 2) revealed a weak latitudinal pattern; species occurring at higher absolute latitudes are smaller, on average, than those at low latitudes. This relationship is significant but has almost no explanatory value ( $r^2 = 0.02$ ; ES4, Table S4.1).

Fig. 2 shows raw data relationships of body length with latitude and the 388 389 environmental predictors; Fig. 3 highlights the main characteristics of univariate 390 correlations (i.e., model coefficients and explained variance; further test statistics in ES4, Table S4.1). Across all models, explained variance is low (pseudo- $r^2 < 0.2$ ) if no or only 391 392 higher-level phylogenetic classifications are integrated into models, while adding genus-393 level classification (hierarchically within higher-level classifications) yields pseudo-r<sup>2</sup> 394 >0.7. Integrating phylogenetic information weakens all fitted environmental effects, but it 395 does not affect the rank order of their strengths, their sign, or their significance (all p 396 <0.001) except for mean annual temperature (p <0.05 when genus-level phylogeny is 397 included).

398 Univariate correlations best support the resource availability hypothesis (positive 399 correlation of body length and NPP). More weakly, data were also compatible with the 400 'converse Bergmann-hypothesis' (positive correlation of body length and mean 401 temperature; Fig. 3) and the seasonality hypothesis (negative correlation of seasonality 402 and size). Univariate correlations are inconsistent in sign with predictions from the TSR 403 and Bergmann rule (both expecting a negative link with mean temperature), and the 404 starvation hypothesis (expecting positive links with temperature extremes).

405 Multivariate modelling (Fig. 4; details in ES4, Table S4.2), as in the univariate 406 models, features low pseudo-r<sup>2</sup>'s unless genus-level classifications are integrated into the 407 models. They also support the link of body length with NPP (i.e., resource availability 408 hypothesis) whereas the positive link with temperature (converse Bergmann hypothesis) 409 is weaker, in particular when phylogenetic data are included. Seasonality effects are weak 410 and inconsistent depending on how much phylogenetic information is included. As in the 411 univariate analyses, multivariate models indicated larger, not smaller species in forested

- 412 habitat. This pattern is not because forests feature higher NPP (which is accounted for in
- 413 models). Unlike with the univariate models, temperature extremes show positive
- 414 coefficients with body length in multivariate models (as expected by the starvation
- 415 hypothesis), but links are weak, non-significant, and VIF analysis indicated predictor416 collinearity issues.
- 417 Based on VIF we simplified this full multivariate model by first dropping 418 seasonality, then temperature extremes. In a final step we also dropped mean annual 419 temperature although it had a very low VIF value because univariate analyses had 420 suggested that it was non-significant and yielded higher AIC values than a model 421 consisting only of forest cover and NPP (ES4, Table S4.2). Finally, a comparison of NPP 422 and forest cover showed that they truly express quite independent aspects of the environment (correlation:  $r^2 = 0.26$ ). Notably, multivariate models do not explain 423 424 substantially more variance overall than univariate models if phylogeny is fully included. 425 Models using other body size metrics as responses, whether univariate or multivariate, 426 lead to the same conclusions (ES5).
- 427

## 428 **DISCUSSION**

429 For 950 sphingid species, occurring from the northern temperate through the 430 African and Asian tropics to the southern temperate zone, we observed distinct 431 geographical patterns of average body size (Fig. 1) that were only weakly related to 432 absolute latitude. Rather, they featured larger species in the Middle East through Southand Southeast-Asia to Melanesia, and smaller species elsewhere, including the African 433 434 tropics. However, species richness in the Middle East is low (Ballesteros-Mejia et al., 435 2017), which increases chance effects (i.e., very high values on the Arabian Peninsula are 436 based on few species per pixel). A comparative, species-focused analysis revealed 437 landscape productivity (NPP) as the strongest correlate of body size, supporting the 438 'resource availability' hypothesis. Other hypotheses of environmental control of body size were weakly (i.e., 'converse Bergmann'; seasonality) or not at all supported; for 439 some hypotheses (i.e., TSR, starvation) even the sign of the expected relationship was not 440 441 met by data. These conclusions held for univariate and multivariate modelling 442 approaches, with or without the inclusion of phylogenetic relatedness of taxa.

However, broad environmental conditions, even NPP, generally explained
relatively little of the variability of body sizes (i.e., <10%) whereas hierarchically</li>
including phylogeny to genus-level consistently produced sound predictions of the global
variability in sizes (i.e., >70% of size variability explained). This implies that
phylogenetic inertia in body sizes, combined with the unknown, past evolutionary events
that shaped the sizes of today's higher-level taxonomic groups' ancestors are much more
relevant for predicting observed body size distributions than current environments.

In light of this it is not surprising that, consistent with other invertebrate studies
(Shelomi, 2012), the much-discussed latitudinal gradient of (average) body sizes,
although statistically observable in data (Fig. 2), is buried under random noise and
explains very little of the data variability (Table S3.1). Because this scatter is wider with
higher species richness towards the tropics (Fig. 2), maximum body sizes feature a much
stronger latitudinal pattern, which shapes the intuitive perception of the pattern (Fig. 1).

456

457 Resource availability effects

458 Our data best supported the idea that body size increases with food availability, 459 which we approximated with productivity (NPP; Blackburn *et al.*, 1999). However, a 460 mechanism whereby plant productivity constrains consumers' body size appears in 461 conflict with the idea of 'energy equivalence' (Damuth, 2007; Brown et al., 2004), which 462 postulates that the variability of species richness, not body size, offsets variability in 463 energy availability. In light of these theoretical uncertainties, we caution against 464 interpreting our statistical support, or that of other studies, as unequivocal proof for the 465 postulated mechanism of the food availability hypothesis. However, we see additional 466 post-hoc support for interspecific food limitation effects on body size in the positive 467 correlation of forest and size. We had considered forests in preliminary analyses to test 468 the agility hypothesis (dense forest benefitting small, agile species in large mammals; 469 Bro-Jørgensen, 2008), but removed this idea from our presentation as quite unreasonable 470 for flying insects (data rejected it comprehensively, not shown ). Although forest data appear statistically independent of NPP (Table S3.2), the recovered positive correlation 471 472 may be due to the fact that a sizable portion of NPP in some regions derives from 473 grassland habitats. However, sphingid caterpillars are folivorous and only a single genus, 474 Leucophlebia, is confirmed as feeding on Poaceae (Diehl, [1982]). The absence of forests 475 may therefore pinpoint where there is little food for sphingids even if NPP may not be 476 particularly low (i.e., in grasslands).

477

#### 478 Other hypotheses

479 We found only weak coefficients for the 'converse Bergmann' and seasonality 480 hypotheses, and multivariate analyses suggest predictor collinearity issues or other 481 artefacts. Recently, Zeuss et al. (2017) presented similar data on geographic patterns of 482 size in European Lepidoptera (and Odonata). While our study went beyond Zeuss et al. 483 (2017) in some important aspects (i.e., larger geographic coverage, more fully including 484 global environmental gradients; comparative analyses accounting for phylogenetic 485 effects), they had data for, and highlighted, a key variable to at least potentially 486 disentangle some mechanistic effects, i.e., voltinism (the number of generations per 487 year). Several hypothetical mechanisms assume, naively, a constant and consistent 488 number of generations per year, which is known to be untrue for many well-studied 489 insect taxa (Zeuss et al., 2017; and references therein). If longer favorable growing 490 seasons or warmer temperatures lead to more generations, this may offset any potential 491 body size increases as postulated by 'converse Bergmann' or seasonality hypotheses.

492 For the other tested hypotheses (i.e., Bergmann/TSR, agility) we did not even 493 observe correlations of the correct sign, which clearly rules them out within the following 494 limitations. The starvation hypothesis postulates that larger species have a lower risk of 495 death due to unexpected climate events, such as starvation or desiccation (Chown & 496 Gaston, 2010). We did not find any support for this idea from negative temperature 497 extremes (which would limit movement and hence foraging), while we could not test, due 498 to data limitations, effects of unusual drought events. However, most growth in sphingid 499 larvae happens during the last instar (e.g., 90% in Manduca sexta; D'Amico et al., 2001), 500 so adult body sizes may not provide a suitable test of postulated starvation effects on the 501 size of earlier instars (i.e., phenology may be more relevant than final size). However, in 502 other insects there has also been little support for this hypothesis except for ants, which 503 are a special case because of their eusocial behavior (Kaspari & Vargo, 1995). Support 504 for the TSR in the literature is mainly from intraspecific studies (including a sphingid; 505 Davidowitz & Nijhout, 2004), but its role is apparently small when it comes to explaining 506 interspecific patterns.

507 Our results are, at least in parts, in conflict with some recent, more localized 508 studies on the body sizes of Lepidoptera and other taxa. Brehm et al. (2019) reported a 509 size increase with elevation in Costa Rica moths, which was best explained by a negative 510 effect of temperature (cf. Bergmann, TSR) but not by an effect of productivity. Beck et 511 al. (2016) also found increasing moth sizes with higher elevations, although they could 512 not link this to flight-time temperatures in their highly seasonal study region, the Swiss 513 Alps. While inconsistent results from ectotherm body size patterns and their causes are 514 common in the scientific literature so far (e.g., Entling *et al.*, 2010; Fattorini *et al.*, 2013; Slavenko & Meiri, 2015; Zeuss et al., 2017; Pallares et al., 2019), it is particularly 515 516 surprising that sphingids, well-known for their (partial) thermoregulation as adults 517 (Heinrich, 1993), did not follow a pattern interpretable towards Bergmann's rule and 518 heat-preservation whereas other moths groups did show such trends in elevational 519 gradient studies. We cannot assess whether methodological differences, peculiarities of 520 elevational studies, a generally high potential for spurious results when dealing with 521 small effect sizes, or other factors may have caused these discrepancies.

522

523 Limitations of the study

524 This study tested, and rejected, some of the global hypotheses regarding 525 environment-body size relationships, based on currently available data on adult sphingid moth sizes. However, our database is necessarily imperfect, and we need to briefly 526 527 discuss how this may have biased the conclusions of our analyses. First, we used an up-528 to-date, but nevertheless coarse and incomplete taxonomic classification as a proxy for a 529 complete, time-calibrated phylogeny with branch length data. Our step-wise inclusion of 530 lower-taxon information gradually increased the explanatory value of statistical models 531 while reducing the tested environmental effects (Fig. 3). We therefore assume that the 532 predominance of phylogeny, as opposed to current environment, in explaining size 533 patterns would be even more pronounced with a more detailed, finely-resolved 534 phylogeny. Second, we only had adult size data available, but many of the mechanistic 535 explanations proposed may apply more strongly to the larval stage, where growth occurs. 536 Thus, concluding that a given hypothesis does not explain observed adult sizes remains valid, but this does not rule out that the proposed mechanisms might play some role in 537 538 shaping the life histories of juvenile stages, which can differ in important aspects (among 539 them, in sphingids, thermoregulation; Heinrich, 1993; Kingsolver et al., 2015). Third, 540 ignoring intraspecific size variation essentially means that we had to view our 541 measurements as a (small) statistical sample around an unknown per-species mean. We 542 do not see any obvious directional bias in this, but it implies that our data may contain 543 considerable random noise simply due to occasionally measuring unusually large or small 544 specimens of a species. As a consequence, the tested effects may be underestimated to an 545 unknown degree. However, the relatively large sample size (i.e., 950 species) would help 546 to counter such chance effects. Programs of computer-aided photography and 547 measurement of specimens in museum collections are under way but they may not 548 directly provide reliable data for solving this issue (Johnson et al., 2013; Trueman & 549 Yeates, 2015). Fourth, our analytical setup involved two potentially confounding effects, 550 phylogenetic inertia and spatial autocorrelation. It is exceedingly complex to control 551 analyses for both effects (Freckleton & Jetz, 2009), and we choose to account in our models for the more important of the two, phylogeny. Strong phylogenetic effects on 552 553 body size have been repeatedly reported in the literature, and they were shown on our 554 data. Spatial autocorrelation occurs in most geographic data and has the potential to 555 affect significance assessments and possibly also coefficient estimates (Bini et al., 2010). 556 However, while we do not wish to give the impression of taking this lightly, it is our 557 assessment that statistically strong patterns rarely lead to changing conclusions when

- applying spatially explicit modelling, whereas the phylogenetic inertia of body sizes most
- 559 probably has the potential to affect conclusions in empirical studies (Diniz-Filho &
- 560 Torres, 2002).
- 561
- 562 Conclusions

563 Average sphingid moth body sizes show distinctive geographic patterns, but they 564 vary only very weakly with absolute latitude. However, maximum body sizes per cell, 565 which are affected by species richness, indicate an increase towards the tropics, which 566 creates the impression of an overall size increase towards the tropics. Among various 567 hypotheses regarding how the environment is shaping such patterns globally, we found 568 strongest support for the food availability hypothesis, as sizes increase with net primary 569 productivity. There was no support at all (i.e., opposite sign of effect) for Bergmann's 570 rule or the temperature size rule, the agility hypothesis, or the starvation hypothesis, 571 while support for the seasonality hypothesis or the converse Bergmann pattern was weak 572 and potentially unreliable. However, phylogenetic effects were much more relevant than 573 any of the tested environmental factors in shaping the observed size data. This implies 574 that past environmental factors in the regions of occurrence of today's higher taxon's 575 ancestors may have shaped observed size patterns, which will be exceedingly challenging 576 to test.

577

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# 737 Data Accessibility Statement

- As electronic supplements to this article we publish our data on sphingid body sizes
- 739 (ES1), a taxonomic classification of species (ES2) as used in analysis, and GIS-
- compatible data of Fig. 1 (ES6). Environmental data used in our analysis are from the
- 741 public domain. Raw distribution maps of sphingid species can be accessed at Map of 742 Life youw molecre
- 742 Life, <u>www.mol.org</u>. \*
- \*) currently only for browsing, by the time of acceptance of this paper they will be
  available for download in GIS format.
- 745

## 746 Appendices: Electronic Supplements

- 747 **ES1** Body size data and specimen sources (spreadsheet in csv-format)
- 748 ES2 Taxonomic classification as used for analyses
- 749 ES3 Supplementary maps and figures
- 750 ES4 Supplementary model statistics
- 751 ES5 Model output for other body size metrics
- 752 ES6 Body size distribution maps (median, interquartile, maxima; ASCI-format)

753

#### 754 FIGURE CAPTIONS

- 755
- **Figure 1** Geographical pattern of the median body length (A), its interquartile range (B)
- and maximum body length found in each cell (C). Grid cells with  $\leq 5$  species were
- removed (map resolution: 100 x 100 km, Mollweide World geographical projection). See
- ES3 for additional maps; data for Fig. 1 are available in GIS-compatible format (ES6).
- 760
- Figure 2 Body length plotted against all predictor variables. LOESS (locally weighted
   scatterplot smoothing) is fitted to indicate main data trends.
- 763

Figure 3 Summary of univariate model results (response: body length) with stepwise hierarchical inclusion of phylogenetic information. Bars (right y-axis) indicate the variance explained by the models, line plots (left axis) the fitted coefficients for the different predictors. Because data are standardized, coefficients can be compared across models as a measure of effect size.

- 769
- Figure 4 Summary of multivariate model results (response: body length) with stepwise
- hierarchical inclusion of phylogenetic information. Bars (right y-axis) indicate the overall
- variance explained by the multivariate model models, line plots (left axis) the fitted
- coefficients for the different predictors.