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Interrelations of global macroecological patterns in wing and thorax size, sexual size dimorphism, and range size of the Drosophilidae

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Abstract: Support for macroecological rules in insects is mixed, with potential confounding interrelations between patterns rarely studied. We here investigate global patterns in body and wing size, sexual size dimorphism and range size in common fruit flies (Diptera: Drosophilidae) and explore potential interrelations and the predictive power of Allen's, Bergmann's, Rensch's and Rapoport's rules. We found that thorax length (r2 = 0.05) and wing size (r2 = 0.09) increased with latitude, supporting Bergmann's rule. Contrary to patterns often found in endothermic vertebrates, relative wing size increased towards the poles (r2 = 0.12), a pattern against Allen's rule, which we attribute to selection for increased flight capacity in the cold. Sexual size dimorphism decreased with size, evincing Rensch's rule across the family (r2 = 0.14). Yet, this pattern was largely driven by the virilis – repleta radiation and only weakly in other lineages. Finally, range size did not correlate with latitude, although a positive relationship with latitude was present in a subset of the species investigated, providing no convincing evidence for Rapoport's rule. We further found little support for confounding interrelations between body size, wing loading and range size in this taxon. Nevertheless, we demonstrate that studying several traits simultaneously at minimum permits better interpretation in case of multiple, potentially conflicting trends or hypotheses concerning the macroecology of insects.

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1 Interrelations of global macroecological patterns in wing and thorax size, sexual

2 size dimorphism, and range size of the Drosophilidae

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17 ABSTRACT

Support for macroecological rules in insects is mixed, with potential confounding 18 19 interrelations between patterns rarely studied. We here investigate global patterns in body and 20 wing size, sexual size dimorphism and range size in common fruit flies (Diptera: 21 Drosophilidae) and explore potential interrelations and the predictive power of Allen's, 22 Bergmann's, Rensch's and Rapoport's rules. We found that thorax length ($r^2 = 0.05$) and wing 23 size ($r^2 = 0.09$) increased with latitude, supporting Bergmann's rule. Contrary to patterns often found in endothermic vertebrates, relative wing size increased towards the poles ($r^2 = 0.12$), a 24 25 pattern against Allen's rule, which we attribute to selection for increased flight capacity in the 26 cold. Sexual size dimorphism decreased with size, evincing Rensch's rule across the family (r² = 0.14). Yet, this pattern was largely driven by the *virilis – repleta* radiation and only weakly in 27 28 other lineages. Finally, range size did not correlate with latitude, although a positive 29 relationship with latitude was present in a subset of the species investigated, providing no 30 convincing evidence for Rapoport's rule. We further found little support for confounding 31 interrelations between body size, wing loading and range size in this taxon. Nevertheless, we 32 demonstrate that studying several traits simultaneously at minimum permits better 33 interpretation in case of multiple, potentially conflicting trends or hypotheses concerning the 34 macroecology of insects.

35 INTRODUCTION

Convergent patterns of phenotypic variation across large-scale environmental gradients have long been recognized and have given rise to several macroecological "rules" predicting such variation as putative adaptive responses to selection. These patterns are generally supported by empirical evidence and underlie theory. Until a more complete understanding of the selective mechanisms underlying the patterns is achieved, however, the predictive power of such rules must remain limited. This is particularly true for insects, in contrast to mammals and birds (Blanckenhorn and Demont 2004; Chown and Gaston 2010; Shelomi 2012).

43 As in all organisms, insect body size is inherently linked to physiology, metabolic rate, 44 survival and reproductive success and is thus thought to evolve in predicted ways if subjected 45 to similar selective drivers (Blanckenhorn 2000, Chown and Gaston 2010). Large size typically 46 increases mating success in males and fecundity in females, but can entail heightened 47 mortality risks and reproductive costs due to prolonged larval development (Blanckenhorn 2000). Amongst the macroecological rules proposed to account for body size variation, 48 49 Bergmann's rule, signifying an increase in size with latitude, is well supported in 50 homoeothermic vertebrates (Bergmann 1847; Meiri and Dayan 2003), but its absence and often 51 its converse is prominent among invertebrates (Shelomi 2012). An increase in size with latitude 52 has been attributed to temperature-dependent variation in growth and metabolic rates, a pattern congruent with the so-called temperature-size rule (Atkinson 1994; Atkinson & Sibly 53 54 1997) that generally predicts insects to grow bigger in the cold (Kingsolver & Huey 2008). 55 However, a shortened active season with increasing latitude can also cause adaptive negative size clines through selection for fast development if development cannot be extended across 56 one season, both within and between species (Chown et al. 1999, Blanckenhorn and Demont 57 58 2004; Zeuss et al. 2017). Small insects with rapid development are thus expected to follow 59 Bergmann clines, whereas large insects may be limited in their development by season length 60 at high latitudes, thus emerging smaller and showing converse Bergmann clines 61 (Blanckenhorn and Demont 2004, Zeuss et al. 2017).

62 In contrast to body size, the relative size of appendages has been found to decrease with 63 latitude in endothermic vertebrates (e.g., Nudds and Oswald 2007, Symonds et al. 2010). Termed Allen's rule, this pattern has received considerable attention and has been attributed 64 to selection for a reduced surface-to-volume ratio to limit heat loss in the cold. Originally 65 66 documented for warm-blooded animals (Allen 1877), qualitatively similar patterns have also been described in invertebrates (e.g. Alpatov 1929, Ray 1960). In insects, most appendages, 67 68 including wings and antennae, are connected to the circulatory system (Chapman et al. 2013), 69 and the constant flow of haemolymph through these appendages can contribute to 70 thermoregulation, as in endotherms. Patterns equivalent to Allen's rule might thus be 71 expected. Still, particularly for small insects, body temperature is unlikely to be strongly 72 dependent on the relative surface area as an insect's body adjusts nearly instantly to the 73 ambient temperature (Harrison and Roberts 2000), although many insect taxa are capable of 74 considerable thermoregulation (social Hymenoptera in particular; e.g., Stabentheiner et al. 75 2010; Chapman et al. 2013). Consequently, small insects primarily regulate their body 76 temperature by modifying their behavior (Clench 1966, Dillon et al. 2009), thus making best 77 use of available microhabitats. The capacity to disperse, however, is greatly restricted at cool 78 temperatures, which impedes take-off in winged insects (Dillon and Frazier 2006, Frazier et al. 79 2008). As increased wing size relative to body size facilitates take-off at cooler temperatures 80 (Frazier et al. 2008), a latitudinal increase of relative wing size (a pattern counter to Allen's 81 rule) can be predicted. Such patterns have indeed been observed in insects (e.g. clinal 82 population differentiation in D. melanogaster: Azevedo et al. 1998), but the repeatability of such

clines awaits further scrutiny. Whether the relative size of insect appendages increases,
decreases or shows any consistent latitudinal pattern at all thus remains unclear.

Whereas both Bergmann's and Allen's rules describe spatial variation in body and 85 86 appendage size irrespective of sex, the widely-studied Rensch's rule focuses on variation in 87 sexual size dimorphism (SSD; Abouheif and Fairbairn 1997; Fairbairn 1997). Rensch (1950) 88 documented that, among closely related species of many disparate taxonomic groups, SSD increases with body size in species in which males are the larger sex but decreases when 89 90 females are larger than males. Unlike Bergmann's and Allen's rules, which capture size 91 variation that likely results from natural selection, differential variation between the sexes has 92 been attributed to a combination of sexual selection on males (e.g., via male-male contest 93 competition) and fecundity selection on females (Fairbairn and Preziosi 1994, Fairbairn 1997, 94 Székely et al. 2004). However, evidence for Rensch's rule is mixed, particularly in taxa with 95 female-biased size dimorphism, such as most insects and spiders (for which SSD tends to 96 increase with body size: Blanckenhorn et al. 2007b; Webb and Freckleton 2007; Stuart-Fox 97 2009), undermining its predictive power.

98 Finally, in addition to body size, populations or species have also been documented to vary 99 in their range size. Specifically, Rapoport's rule (or "effect": Stevens 1992) predicts that species 100 occurring close to the equator will have more restricted ranges because they exhibit limited 101 climatic tolerance as a result of adapting to a local environment with low climatic variability. 102 Conversely, species occurring at higher latitudes (or altitudes), which feature ample climatic 103 variability, are predicted to be better adapted to colonize and occupy more diverse habitats. 104 Evidence for this rule is still controversial as it might be restricted to specific latitudes and 105 certain regions and somewhat scale dependent (Rohde 1996, Ruggiero and Werenkraut 2007). 106 The four macroecological patterns described above are often studied in isolation from each 107 other. Nevertheless, body size, wing morphology (which relates to dispersal capacity; e.g., 108 Ray et al. 2016), SSD and range size are all likely to be ecologically and evolutionarily 109 interrelated, if only because most adaptive explanations discussed above relate to climate. For 110 example, although Rensch's rule has mostly been studied independently of environmental 111 factors, latitudinal patterns of SSD have been observed (Blanckenhorn et al. 2006), and a 112 relationship of Rensch's rule with the widely-observed temperature-size rule was postulated 113 but not found after all (Hirst et al. 2015). Similarly, a large body of literature investigates 114 relationships between range size and both dispersal capacity and body size (Gaston and 115 Blackburn 1996, Malmqvist 2000, Lester et al. 2007, Rundle et al. 2007, Laube et al. 2013), thus 116 raising the issue of whether clinal variation in size and dispersal might generally drive 117 Rapoport's rule, possibly undermining its ecological relevance (Reed 2003). It is thus useful -118 if not imperative - to account for additional potentially confounding effects when investigating 119 range-size variation (Laube et al. 2013).

120 Rigorous examination of macroecological patterns requires not only robust taxon sampling 121 and coverage of a large geographic range, but also the ability to account for phylogenetic 122 relationships among species. The Drosophilidae, a highly speciose and diverse family of fruit 123 flies with a global distribution, should be highly suited to assess such patterns and their 124 underlying mechanisms in insects. Many drosophilids are cosmopolitan generalists, whereas 125 others are highly specialized and endemic to small geographic areas (e.g., see Ashburner 1981). 126 In this taxon, Bergmann's rule has received considerable attention at the intraspecific level 127 along both latitudinal and altitudinal gradients (e.g., D. melanogaster: van't Land et al. 1999, 128 Fabian et al. 2015, Klepsatel et al. 2014; D. buzzatii: Karan et al. 2000; D. subobscura: Gilchrist et 129 al. 2001; Zaprionus indianus: Karan et al. 2000), but little information is available at the

interspecific level. Similarly, Rensch's rule has been addressed in the *obscura* group (Huey et
al. 2006), but the general pattern has not been investigated (but see Blanckenhorn et al.
2007a,b). The potential relationship of range size with body and appendage size also remains
largely unexplored in this group.

134 We here assessed in a comparative framework the ecogeographical patterns of thorax 135 length and wing size, SSD, and range size in 151 drosophilid species from around the globe, 136 including members of all three major *Drosophila* clades plus species of other genera belonging 137 to this speciose family. Accounting for phylogenetic non-independence based on a 138 reconstructed phylogeny, we investigated the geographical patterns described above and 139 potential relationships between the four macroecological rules. Such relationships are 140 expected under the prediction that they all relate (to some extent) to climate, but their 141 covariation is poorly studied empirically. We thus aimed to better understand the causes and 142 consequences of macroecological variation.

143 MATERIALS & METHODS

144 **Phylogeny reconstruction**

145 To reconstruct the drosophilid phylogeny, we obtained the sequences of six nuclear, three 146 mitochondrial and three mitochondrial ribosome genes from GenBank (see Supplementary 147 Table S1 for accession numbers and sequence coverage). The gene coverage per species ranged 148 between 1 and 12 (mean \pm SD = 7.2 \pm 2.9 loci/species), with a total sequence length of 6,269.2 \pm 149 3,267.6 bp (range = 337 – 14,449 bp). The nuclear sequences comprised the genes expressing 150 the alcohol dehydrogenase (Adh), α -amylase-related protein (Amyrel), aromatic-L-aminoacid decarboxylase (DOPA decarboxylase; Ddc), glycerol-3-phosphate dehydrogenase 151 152 (Gpdh), and xanthine dehydrogenase (Xdh). The mitochondrial genes included the 153 cytochrome c oxidase subunits I, II, and III (COI, COII and COIII, respectively) and the NADH 154 dehydrogenase subunit 2 (ND2). The ribosomal genes included 28S and the large and small 155 subunits of 12S and 16S (omitting the adjacent tRNAs as they were difficult to align and 156 represented only a small amount of data). For each locus, we aligned the sequences of all species using multiple sequence alignment (MUSCLE) as implemented in MEGA v.7.0 (Kumar 157 158 et al. 2016) and determined its best nucleotide substitution model using jModelTest v.2.1.7 159 (Darriba et al. 2012). The best substitution models were GTR+Γ+I for all nuclear genes and 16S, 160 HKY+Γ+I for the mitochondrial genes and 12S, and HKY for 28S, respectively.

161 Subsequently, we reconstructed the phylogeny based on Bayesian inference using BEAUTi 162 and BEAST v.1.8.3 (Drummond et al. 2012), with unlinked substitution models, a relaxed 163 uncorrelated log-normal clock, and a Yule speciation process. Due to a lack of well-defined 164 fossil dates in our sample of species, and because the absolute timing of speciation events was 165 deemed less important for our analyses than the relative branch lengths, we omitted the time 166 calibration. We ran the Markov Chain Monte Carlo (MCMC) simulation on the CIPRES Science 167 Gateway (http://www.phylo.org; Miller et al. 2010) for 100 million generations, sampling 168 every 10,000th tree. We used Tracer v.1.6 (Rambaut and Drummond 2013) to examine the 169 convergence of the Bayesian chain and the stationary states of all parameters, considering 170 effective sample sizes (ESSs) greater than 200 to be adequate. Finally, we generated a 171 maximum clade credibility tree with mean node heights and a 10% burn-in using 172 TreeAnnotator v.1.8.3 (Drummond et al. 2012).

173 In addition to the full *Drosophila* phylogeny, we separately generated a phylogeny for the 174 *Zaprionus* dataset (for which only body lengths and no thorax lengths were available). For 175 these species, adequate sequence coverage was restricted to the Amyrel, COI, COII, and 28S genes, respectively (see Supplementary Table S2). We used the same procedures as above,
except that GTR+Γ+I was the best substitution model for all loci and the MCMC chain was run
for only 30 million generations, with a tree sampled every 3,000 generations.

179 Data collection

We measured sex-specific body size as thorax length of field-caught specimens (distance between the tip of the scutellum and the basis of the head, a standard measure) for 56 species of Drosophilidae stored at the Zoological Museum of the University of Zurich. Whenever available, we measured at least 10 individuals per sex per species. We further obtained thorax length data for 111 additional species and data on total body length for 20 *Zaprionus* species from the literature (see Supplementary Table S3).

186 For each of 146 database TaxoDros our species available in the 187 (http://www.taxodros.uzh.ch/), we retrieved the geographic coordinates of every faunistic 188 record. This database comprises an enormous amount of information on the taxonomy and 189 distribution of more than 6,800 species of drosophilids. All coordinates are derived from 190 published data or stem from museum catalogs in which the identification was verified by 191 experts. Although these faunistic data do not stem from standardized collecting schemes but 192 are somewhat haphazard in nature, TaxoDros represents one of the most powerful and 193 comprehensive data sources for any insect taxon. Nonetheless, the sampling coverage might 194 be biased towards certain regions and not be homogeneous across the globe, a common 195 problem when handling such datasets. Thus, we concede that any inference based on these 196 distribution data should be treated with some caution (see Conclusion).

We included only species with at least 20 unique sampling locations in our analyses(removing duplicate and nonsensical localities, resulting in over 25,000 unique coordinates;

mean number of coordinates per species: 273.3, SE: 47.9, median: 137; Fig. 1). Many drosophilids are distributed globally, but their range is often restricted within latitudinal bands such that they do not occur at the equator. We thus used the mean of the *absolute* latitudinal distribution to obtain a suitable estimate of the species-specific latitudinal distribution.

204 Although several major ecogeographic rules describe patterns of latitudinal trait variation, 205 latitude itself remains a compound trait integrating various climatic factors. In this respect, the 206 differential effects of temperature and seasonality are of particular interest. Using climatic 207 data, we tried to decompose latitude into variates related to temperature or seasonality. 208 However, due to the high collinearity among climate variables and latitude, we were unable 209 to use multiple-regression approaches (variance inflation factor always greater than 5 and 210 often greater than 10). When using a principal component analysis (with oblique rotation) to 211 extract the major axes of variation, both temperature and seasonality variables loaded strongly 212 on the first principal component (but in opposite directions, thus mirroring latitude), while all 213 other dimensions did not explain significant proportions of variance and did not show any 214 association to either temperature nor seasonality (Supplementary Table 1). For the sake of 215 interpretation (as we were unable to disentangle temperature from seasonality and latitude), 216 we decided to restrict our analyses to latitudinal patterns alone, although future studies 217 disentangling the climate compounds of latitude are clearly desirable.

218 Thorax length

We analyzed the relationship between log thorax length (mean of male and female values) and median latitude, using phylogenetic generalized linear models (PGLS) as implemented in the R package *caper* (Orme et al. 2012). We used the sexual dimorphism index (SDI) as an estimate of the strength and direction of SSD as proposed by Lovich and Gibbons (1992). We thus divided the thorax length of the larger sex (usually the female) by the smaller and subtracted 1 from this ratio, which arbitrarily defines the SDI positive if females are the larger sex and negative if males are larger.

Wing size

We retrieved wing-size data for 54 species from Bolstad et al. (2015). These wing sizes 227 228 represent the square root of wing area derived from outline spline reconstructions. To assess 229 clinal variation in relative wing size, we used PGLS with latitude as the predictor and thorax 230 length as a covariate. In addition, we calculated wing loading, which is typically associated with wing-beat frequency and flight capacity (Pétavy et al. 1997; Frazier et al. 2008). Wing 231 loading is usually defined as some ratio of body mass and wing area, where low values relate 232 233 to better dispersal capacity as less weight is "loaded" onto the wing. Because body mass 234 estimates were lacking, we used thorax length³, which scales well with mass. We analyzed its 235 relationship with latitude using PGLS.

236 Sexual size dimorphism

To test whether SSD scales iso- or allometrically with body size, we applied phylogenetic reduced major-axis regressions (as implemented in the R package *phytools*: Revell 2012) of log male against log female thorax length across all species (for justification see Fairbairn 1997; Blanckenhorn et al. 2006). Rensch's rule is evident only if the slope of this relationship exceeds one. We repeated the analysis separately for all three major clades of *Drosophila* (*Sophophora* subgenus, *immigrans-tripunctata* radiation, *virilis-repleta* radiation) and the *Zaprionus spp*. data set, for which body size was measured as total body length. To quantify the predictive strength of Rensch's rule in drosophilids, we further calculated r² from a PGLS of SDI against log mean
size.

246 Range size

247 When investigating variation in range size, we considered only species with 20 or more 248 unique records, thus reducing the number of species with sufficient data to 110 (mean number 249 of coordinates per species: 273.3, SE: 47.9; median: 137). To approximate range sizes directly 250 from faunistic records, we derived range-size estimates using α -hulls (as in Gallagher 2016), 251 which are more robust than simple minimum convex polygons, particularly when sampling 252 is haphazard and not standardized (Burgman and Fox 2003). We used the Lambert azimuthal equal-area projection to generate appropriate range-size estimates in km² across the globe and 253 254 restricted these range sizes to actual land masses using the R package rangeBuilder (Davis 255 Rabosky et al. 2016). A PGLS model was used to test for a relationship between log range sizes 256 and latitude (median). In order to test for potential confounding effects of body size and 257 relative wing length (as an estimate of short-distance dispersal), we also performed multiple 258 PGLS regression analyses with thorax length and wing size as covariates. As wing sizes were 259 available for only 54 species, the sample size for this analysis was drastically reduced (note, 260 however, that the number of sampling coordinates per species was greater in this reduced data 261 set: mean: 395.9, SE: 89.7; median: 176).

262

263 **RESULTS**

264 Thorax length and wing size

Log mean thorax length and wing size increased with median latitude (thorax length: r = 0.23 [95% confidence limits: 0.04, 0.39], $\lambda = 0.97$, n = 107, P = 0.019; wing size: r = 0.30 [0.04, 0.51],

 $\lambda = 0.96$, n = 54, P = 0.026; fig. 1), although the unexplained variation in thorax and wing size was rather large (fig.1). Log wing size also showed a positive relationship with latitude when log thorax length was included as a covariate (r = 0.30 [0.03, 0.50], λ = 0.55, n = 54, P = 0.031), suggesting a disproportionate increase in wing size towards high latitudes. Accordingly, wing loading decreased with latitude (r = -0.35 [-0.09, -0.54], λ = 0.00, n = 54, P = 0.009; fig. 1).

272 Sexual size dimorphism

273 Phylogenetic signals in male thorax length ($\lambda = 0.86$, P < 0.001), female thorax length ($\lambda =$ 0.84, P < 0.001), mean body size (λ = 0.85, P < 0.001) and SSD (λ = 0.75, P < 0.001, fig. 2) suggest 274 275 phylogenetic inertia of body size and SSD in drosophilids (fig. 2). When testing Rensch's rule across all species, RMA slopes were significantly steeper than unity (β_{phyIRMA}: 1.10 [1.06, 1.15], 276 277 P < 0.001, n = 151; fig. 3), and body size explained 14% of the total variation in SDI. RMA slopes 278 did not significantly differ between radiations (log female size x radiation interaction: $F_{2,114}$ = 279 2.31, P = 0.104), however, when testing Rensch's rule within the three major radiations, we did 280 not find consistent support. Rensch's rule was evident in the *virilis - repleta* radiation (β_{phyIRMA}: 281 1.08 [1.01, 1.14], P = 0.021, n = 48; fig. 3), but not in the *immigrans – tripunctata* radiation ($\beta_{phyIRMA}$: 282 1.02 [0.92, 1.13], P = 0.650, n = 22; fig. 3). In the *Sophophora* subgenus, the RMA slopes were very steep ($\beta_{phyIRMA}$: 1.18 [1.06, 1.31], P = 0.002, n = 51; fig. 3), but this pattern was driven exclusively 283 by Drosophila prolongata. This species is by far the largest member of this subgenus and the 284 285 only one showing male-biased SSD. When excluding D. prolongata, Rensch's rule was no longer supported in this clade (β_{phyIRMA}: 0.99 [0.92, 1.08], P = 0.996, n = 50; fig. 3). The 286 287 relationship of male and female body length also did not deviate from isometry in Zaprionus 288 *spp.* (β_{phyIRMA} : 1.06 [0.83, 1.28], P = 0.577, n = 16; fig. 3). There was also no evidence for a 289 correlation between sexual size dimorphism and latitude (r = 0.09, [-0.10, 0.27], λ = 0.67, n = 290 107, P = 0.369).

291 Range size

292 Log range size did not correlate with latitude in the simple linear model using the full data 293 set (r = 0.03 [-0.16, 0.22], λ = 0.03, n = 105, P = 0.777, fig. 4). However, in a phylogenetic multiple 294 regression including thorax and wing size as additional explanatory variables, range size 295 increased towards the poles (r = 0.37 [0.11, 0.56], P = 0.007, λ = 0.98) whereas thorax and wing 296 length had no effect on range size (thorax: r = 0.10 [-0.17, 0.36], P = 0.469, $\lambda = 0.98$; wing size: r 297 = 0.04 [-0.23, 0.30], P = 0.770, λ = 0.98). Note that the data underlying this multiple regression 298 represent only a subset of the data because wing size, thorax length and range extent data were 299 available for only 54 species.

300

301 **DISCUSSION**

302 Our study of the morphology and global distribution patterns of Drosophilidae lends 303 support to several macroecological phenomena. In accordance with Bergmann's rule, thorax 304 length and wing size increased with latitude, and the same was true for relative wing size 305 (contrary to Allen's rule). Our data on SSD also support Rensch's rule overall, but this pattern 306 was mostly driven by the virilis - repleta radiation, with weak support in three other major 307 clades. We found no further evidence for a latitudinal cline in SSD. Range size did not vary 308 with latitude across all species of our study, not generally supporting Rapoport's rule. 309 However, when controlling for the potentially confounding effects of body size and shape 310 (and thus reducing our dataset), we found a significant increase in range size with latitude. In 311 the following, we link our results to the ecology and physiology of drosophilids and discuss potential causes and consequences of these macroecological patterns and their apparent 312 313 idiosyncrasy depending on which species are analyzed.

314 Latitudinal effects on body size, shape and range size evolution

315 Bergmann's rule is thought to be driven by variation in temperature (Atkinson and Sibly 316 1997, Shelomi 2012), whereas its converse represents an adaptive response to season length 317 (Blanckenhorn and Demont 2004). Drosophilids generally follow a weak positive Bergmann 318 cline, thus at best suggesting only minor effects of temperature and no role of season length in 319 this family. As most drosophilids are small, fast-developing and therefore strongly 320 multivoltine (although some univoltine species and populations exist; e.g., Lakovaara et al. 321 2009), this could be expected (Blanckenhorn and Demont 2004). Compared to the strength of 322 interspecific clinal variation in other insects, the variation explained by latitude in wing ($r^2 =$ 323 0.09) and thorax length ($r^2 = 0.05$) is below average, though not particularly low (cf. r^2 for 324 similar interspecific comparisons from Shelomi (2012): mean = 0.22, median = 0.10, SD = 0.25, 325 n = 18). Given that most individuals measured for this study were collected in the field and 326 not raised under controlled environments, a considerable amount of body size variation must 327 be attributable to phenotypic plasticity. Although this typically also applies to other studies of 328 various taxa, it is possible that we underestimate the strength of the latitudinal pattern.

329 Even though processes acting within species (sometimes termed neo-Bergmannian rule or 330 James's rule) do not necessarily coincide with among-species patterns (Blackburn et al. 1999), 331 the interspecific clinal variation observed here is consistent with analogous intraspecific 332 variation in drosophilids (e.g., Chown and Gaston 2010). Due to this qualitative consistency, 333 it is reasonable to assume a common underlying mechanism. However, following the 334 temperature-size rule (Atkinson 1994), drosophilids tend to grow larger in cool environments 335 in general (e.g. Ray 1960), and experimental laboratory rearing would be required to test 336 whether this between-species pattern is driven by evolutionary or purely plastic (i.e. physiological) responses. Note, however, that intra-specific common-garden experiments
suggest a strong genetic component (e.g., James et al. 1995).

339 Along with thorax length, wing size increased with absolute latitude, though its steeper 340 increase resulted in disproportionately larger wings at higher latitudes and consequently 341 lower wing loading. Because log wing length showed an isometric relationship with log thorax 342 length across species (evolutionary allometric coefficient derived from a phylogenetic reduced 343 major axis regression: β = 0.94, P = 0.528), allometric scaling relationships cannot explain the 344 relative increase in wing size with latitude. In contrast to warm-blooded animals, in which 345 latitudinal variation in appendage size has been attributed to selection for thermoregulatory 346 efficiency (e.g. reduced bill size in birds: Symonds et al. 2010), such mechanisms seem unlikely 347 to act in insects. Yet, thermoregulation may still be involved in shaping the observed pattern. 348 Being unable to control body temperature endogenously, small insects such as drosophilids 349 regulate body temperature mostly by modifying their behavior (Dillon et al. 2009, Kjærsgaard 350 et al. 2010). Since flight is hampered in the cold and larger wings lower the temperature 351 threshold for take-off (Dillon and Frazier 2006, Frazier et al. 2008), relatively larger wings near 352 the poles could represent an adaptation to large climatic variability or low temperatures 353 (Angelo and Frank 1984, Pivnick and McNeil 1986, Azevedo et al. 1998, Dillon et al. 2009). Such 354 correlations between dispersal capacity and latitude or altitude have been documented in 355 several species (Hassall 2015, Kjærsgaard et al. 2015, Rohner et al. 2015), including latitudinal 356 clines for wing loading in D. melanogaster (Azevedo et al. 1998, Klepsatel et al. 2014), and again 357 suggest a common underlying mechanism of intra- and interspecific clines. Yet, greater 358 dispersal capacity may be an essential prerequisite for colonizing habitats at high latitudes in 359 the first place (e.g., following colonization after the last glacial period). Invading less 360 predictable habitats may then in turn promote the evolution of physiological adaptations that

361 are linked to the climate experienced. This alternative explanation seems unlikely, however, 362 given that wing loading did not correlate with range size (see below), and therefore a direct 363 link between colonization success and short-range dispersal ability appears questionable. 364 Increased relative wing size could thus indeed be associated with cold temperature or increased climatic variability per se and, consequently, a common pattern in small pterygote 365 366 insects. Whether this pattern constitutes evidence against Allen's rule as originally formulated 367 or whether such patterns should be discussed in this context at all is certainly debatable, but 368 also not the main point here.

369 When considering all data, we found no significant correlation between range size and 370 absolute latitude, suggesting no support for Rapoport's rule overall. Our multivariate analysis 371 further suggests that wing size (a proxy for short-range dispersal potential) and thorax length 372 do not confound this relationship, even though both traits have been shown to play major roles 373 in range-size evolution (Malmqvist 2000, Lester et al. 2007, Rundle et al. 2007, Gaston 2009, 374 Swaegers et al. 2014). Surprisingly, the subset of species for which both wing and thorax data 375 were available showed a significant increase of range size with latitude. This discrepancy is 376 unlikely explained by phylogeny, as the species used in the multivariate analysis are well 377 distributed across all major clades (see fig. S1). At the same time, there was more faunistic 378 information available for these 54 species (based on the number of coordinates), thus likely 379 increasing the precision of our range-size estimation. Alternatively, this deviating pattern 380 could also be caused by the ecology of these particular species. Unlike many other drosophilids 381 (Markow and O'Grady 2006), these species can be cultured easily in the laboratory, so they 382 may be particularly flexible and undemanding in their ecological preferences. If so, their range 383 size might be less affected by ecological barriers such as the distribution of substrate host

384 species. Given this non-random subset and the non-standardized sampling scheme used to385 derive range size, future research should test these patterns at the global scale.

386

Sexual size dimorphism and Rensch's rule

387 Females were the larger sex in nearly all species investigated. There is, however, one 388 particularly conspicuous exception to this trend: Drosophila prolongata. This species is not only 389 the largest-bodied species in the *Sophophora* subgenus, but also the only species exhibiting 390 pronounced male-biased sexual size dimorphism (also see Rohner et al. 2017). This species 391 adds further evidence to the notion that there is great potential for rapid evolution of reversed 392 SSD in Diptera, and its apparent association with increased male-male contests (Rohner et al. 393 2016), which are also common in D. prolongata (Kudo et al. 2015). Given its large size and male-394 biased SSD, D. prolongata strongly affected the statistical appraisal of Rensch's rule here (Fig. 395 3), reemphasizing potential issues with the classic assessment of Rensch's rule when male- and 396 female-biased taxa differ in size (see Webb and Freckleton 2007). Nevertheless, the mating 397 system, including the evolutionary drivers of SSD and body size, of *D. prolongata* is likely to 398 be derived, such that this single extraordinary species may obscure rather than testify to 399 Rensch's rule in Sophophora.

Although we found support for Rensch's rule across the entire family, this pattern did not hold within some of the major (sub)radiations. Nevertheless, in most cases the reduced majoraxis slope between males and females was steeper than one. In fact, empirical research demonstrates frequently that support for Rensch's rule depends strongly on the taxonomic level with considerable variation among closely related clades (e.g., Webb and Freckleton 2007). Even if supported in interspecific comparisons, Rensch's rule does not necessarily hold among or within populations of these species (Blanckenhorn et al. 2007a). Within-population 407 variation in SSD is likely driven, at least in part, by ontogenetic processes and thus not 408 necessarily linked to selective forces driving Rensch's rule across species (Teder and Tammaru 409 2005). In theory, Rensch's rule should nonetheless hold across populations and species. Sexual 410 selection on male size tends to be the strongest and most consistent evolutionary driver of 411 large male size (e.g.: Székely et al. 2004, Rohner et al. 2016), and Rensch's rule is arguably more 412 prominently supported in taxa with male-biased SSD due to this selective homogeneity (e.g., 413 Stuart-Fox 2009). In contrast, fecundity selection mediates female-biased SSD to a much lesser 414 extent than expected, as selection for small male size or other evolutionary scenarios are also 415 common (Pincheira-Donoso and Hunt 2015). Blanckenhorn et al. (2007b) and Huey et al. (2006) 416 found evidence for Rensch's rule in Drosophila based on 23 and 42 species, respectively. Our 417 data suggest that this result is robust, but driven mostly by the *virilis - repleta* radiation, while 418 tests in other clades do not support deviations from isometry.

419 **Conclusions**

420 Macroecological rules are sometimes considered to be weak and idiosyncratic, partly 421 because their predictive strength and manifestation varies across taxa, but possibly also 422 because they might be interrelated or confounded. While our comparative analyses largely 423 corroborate previously reported intraspecific patterns for thorax length and wing size, support for Rensch's rule was inconsistent among clades (although slopes did not significantly differ 424 425 between clades). Moreover, we found no support for Rapoport's rule overall and showed that 426 this pattern is not necessarily associated with thorax length or wing size of high-latitude 427 species. Although entirely correlational, we further suggest that increased relative wing size 428 at higher latitudes may be driven by selection for more efficient flight and thermoregulatory 429 behavior.

430 We conclude that studying the relationships between several prominent macroecological patterns can shed more light on broad ecogeographic patterns. However, we here found only 431 432 little evidence for confounding effects. Nevertheless, given that their putative underlying 433 causes are often linked to climatic factors, considering several macroecological patterns 434 simultaneously at minimum permits better interpretation in case of multiple, potentially 435 conflicting trends or hypotheses, as was the case here for wing size and its potential 436 relationship with Allen's rule. Future research should focus on the underlying physiological mechanisms to definitively discern the causes and consequences of various macroecological 437 438 patterns in Drosophilidae and other taxa.

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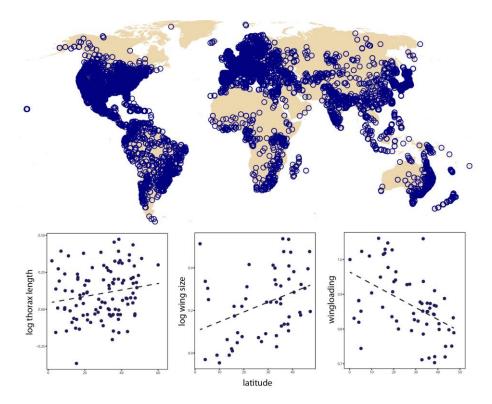
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630 Figure 1: Top: Species specific range size estimates and mean latitudinal distributions were 631 derived from over 25,000 unique sampling locations depicted here. This global dataset was 632 retrieved from TaxoDros, a large and detailed database on taxonomy and diversity of 633 drosophilids (http://www.taxodros.uzh.ch/). Bottom: thorax length and wing size increases 634 with mean absolute latitude, demonstrating a weak interspecific Bergmann cline in drosophilids. Wing size increased more strongly with latitude than thorax length, resulting in 635 lower wing loading (thorax length³/wing area) towards the poles. These plots showing simple 636 637 linear regressions are for illustrative purposes only. All analyses were done using PGLS 638 (phylogenetically corrected correlation coefficients are given in the text).

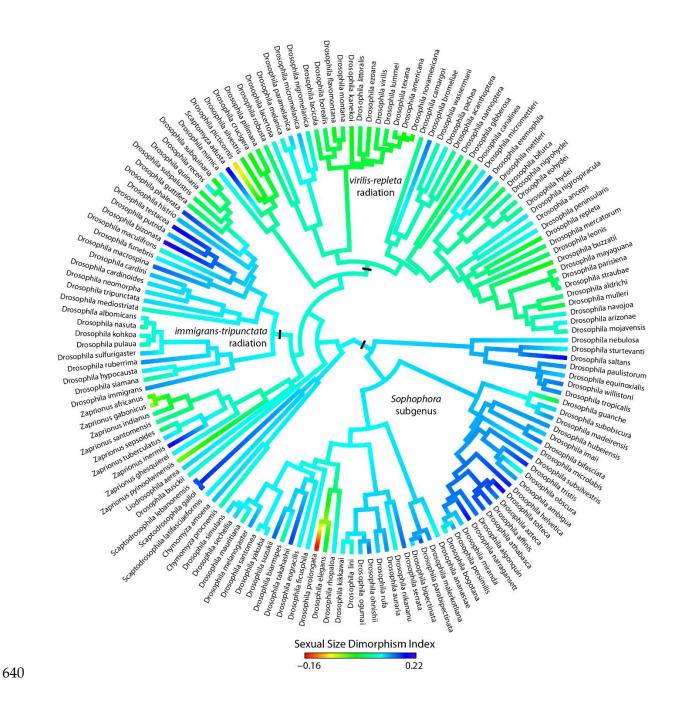


Figure 2: Ancestral state reconstruction of sexual size dimorphism (SDI = (thorax length of larger sex / thorax length of smaller sex) -1; arbitrarily defined negative if males are the larger sex), for illustration purposes only. SSD shows strong phylogenetic signal and in most species females are the larger sex (positive SDI values). The conspicuous exception is *Drosophila prolongata*, which shows pronounced male-biased SSD.

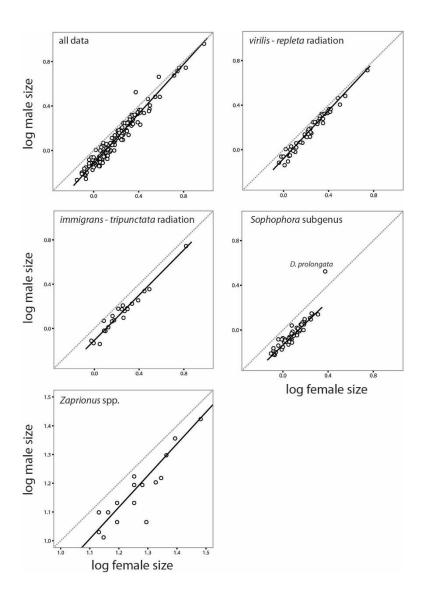




Figure 3: Male size increases more with body size than female size, supporting Rensch's rule for drosophilids. This pattern is however mostly driven by the *virilis-repleta* radiation and absent in all other major clades. For *Zaprionus spp.*, only total body length was available while for all other species thorax length was used. Note that regression lines are derived from nonphylogenetic major axis regressions for illustrative purposes only.

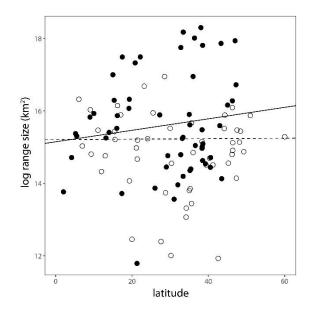


Figure 4: Alpha-hull derived range sizes did not correlate with latitude when all species were analyzed (broken regression line). However, when applying a multiple PGLS regression controlling for thorax length and wing size, range size increased with latitude (solid regression line). Based on merely a limited number of species, this does not seem to be a general pattern across the family, although species for which wing size data were available are dispersed well across the family . The plot shows simple linear regressions for illustrative purposes only. All analyses were done using PGLS (phylogenetically corrected correlation coefficients are given in the text).

670 Supplementary files:

671 Table S1: Principal component analysis of four bioclimatic variables did result in one

672 significant axis of variation. Given that the variables related to temperature load opposite to

673 climate variation variables, the first principal component essentially represents latitude and

674 the effects of temperature and variation cannot be disentangled.

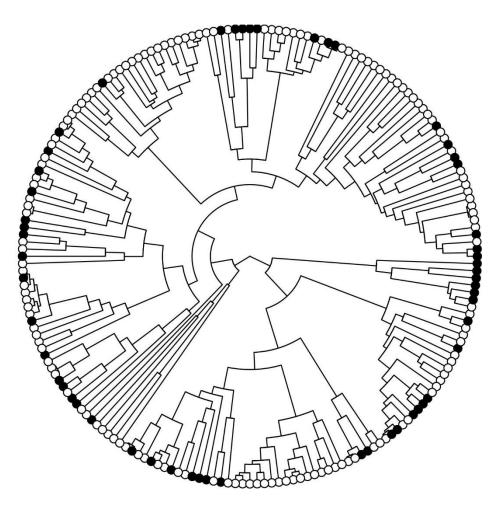
Proportion of variance explained					
	PC1	PC2	PC3	PC4	
Eigenvalue	3.29	0.68	0.03	0.00	
Proportion of variance explained	0.82	0.17	0.01	0.00	
Cumulative variance	0.82	0.99	1.00	1.00	

675

PC loadings			
	P 6 76		
Annual Mean Temperature	-0.98		
Temperature Seasonality	0 ,9 5		
Temperature Annual Range	0.91		
Mean Temperature of Warmest Quarter	-0.78		
	678		

679

Figure S1: Taxa for which distribution data were available are well dispersed across thephylogeny.



○ only distribution data

• distribution and wing size data

689