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## **Abstract**

 Evidence is mounting that migration in ectotherms is more widespread than formerly believed. Thus, a number of insects living in temperate climates, including locusts, butterflies, moths and dragonflies following seasonal migration strategies show high responsiveness for alterations in climatic processes, similarly to a broad taxonomic range of birds. On global scales, migratory insects include iconic large butterflies, dragonflies and also a number of crop pests. However, insect migrations are ecologically distinct from those of vertebrates, often relying heavily on seasonal winds and multiple generations to complete a full annual cycle, due to short insect life spans. Here we review publicly available online resources to identify key patterns of spatial, taxonomic scales and complexity of climatic responsiveness to environmental predictors in migratory insects. We found that migratory insects respond to various levels of complexity in climatic patterns and these responses are predicted by life history and ecological traits: i) responses to climatic effect type was predicted by climate zone(s) of the distribution area, whereas ii) response to climatic complexity was predicted by body size. In conclusion, migratory insects respond to various levels of complexity in climatic processes and this responsiveness is governed by a substantially wider array of environmental predictors than demonstrated in vertebrates.

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### **Introduction**

 During the past decades, an increasing number of investigations revealed that migration - when individuals regularly travel between the breeding and non-breeding areas along directed paths - is a

 key life history trait not only in birds and mammals, but also in ectotherms, such as insects and crabs (Rainey 1963; Schaefer 1969; Urquhart & Urquhart 1978; Dingle 1996). On global scales, migratory insects include iconic large butterflies (e.g. *Danaus plexippus*), dragonflies (e.g. *Pantala flavescens*) and also a number of crop pests (e.g. *Agrostis ypsilon*) (Holland *et al*. 2006; Chapman *et al*. 2015). Moreover, other less conspicuous groups of insects have also been shown to migrate, such as hoverflies (Wotton et al. 2019) and aphids (Bell et al 2015). Insect migrations are ecologically distinct from those of vertebrates, often making use of seasonal winds and relying on multiple generations to complete a full annual cycle, due to short insect life spans (Chapman et al. 2011).

 Seasonal migrations of insects vary widely in their spatial patterns (Satterfield et al. 2020), encompassing both the well-known round-trip migrations of numerous hoverfly, dragonfly, and butterfly populations, including the Monarch Butterfly (*Danaus plexippus*) and the Painted Lady (*Vanessa cardui*) (Chowdhury et al. 2021; Stefanescu et al. 2007, Stefanescu et al. 2013, Hu et al 2021), as well as the more complex movements of Desert Locusts (*Schistocerca gregaria,* Kennedy 1951) and Australian Armyworm Moths (*Mythimna convecta,* Farrow & McDonald 1987), migratory journeys of which form interconnected networks across continents. Other insects like Mustard Aphids (*Lipaphis erysimi*) and Green Lacewings (*Chrysoperla carnea*) demonstrate seasonal migrations radiating out from source sites, resulting in repeated population-scale patterns (Drake & Gatehouse 1995).

 Studies revealed that several insects (e.g. Oleander Hawkmoths [*Daphnis nerii*] in Japan; Ohba *et al.* 1999) are altering the phenology or topological properties of their seasonal movements in response to environmental change. Moreover, some migrants have already shifted their ranges northwards (Satterfield *et al.* 2020) as a response to warmer temperatures (Sparks *et al.* 2005). For example, Juhász *et al.* (2021) found that the relative intensity of the first and last migratory peaks of the Painted Lady significantly increased and showed temporal advancement during the past decade.

 Migratory insects may be especially vulnerable to current global climatic trends, owing to their dependence on spatially distributed resources, which may be differentially influenced by changes in climate (Lemoine & Böhning-Gaese 2003; Oberhauser & Townsend Peterson 2003). For example, the Cloudless Sulphur (*Phoebis sennae*) and Question Mark (*Polygonia interrogationis*) butterflies in the US (Breed *et al.* 2013), as well as the Giant Honeybee (*Apis dorsata*) in India (Sihag 2014) show marked climatic responsiveness mediated by the distribution of their food plants. Similarly, even the once highly abundant Rocky Mountain Locust (*Melanoplus spretus*) has been shown to be driven to the verge of extinction by adverse environmental processes: this locust once performed dry-season migrations in invasion-like numbers from high-altitude to low-altitude zones in the Rocky Mountains (Sorensen 1995). Additionally, once widespread macro-moths (including pests) are also declining in the UK, including the migratory Silver Y (*Autographa gamma*), Black Cutworm (*Agrotis ipsilon*), and Turnip Moths (*Agrotis segetum*) (Fox 2013). Indeed, climate change has been confirmed to modulate the dynamics of migratory insect populations (Fox et al. 2014, Conrad et al 2004).

 A key component of analysing how changes in climate impact the migration strategy of various species is the identification of the most influential climatic parameters. Theory predicts that migratory species will vary in their sensitivity to meteorological conditions and that this responsiveness may be time-dependent and may change over the range of the species (Robinson *et al.* 2005). Further, insects' responses to climatic conditions have been shown to vary on the levels and scales of climatic processes: climatic responsiveness might be dependent on i) causal mediation, defined as direct (i.e. the response is strongly correlated with the climatic parameters) or 94 indirect (i.e. the response is not related directly to climatic processes but strongly correlated with an ecological process or pattern which is governed by climatic trends) effects; ii) spatial scale, defined as local (i.e. confined to a region) or continental (i.e. covers at least 50% of a continent); and iii) complexity, classified as responses to a single or multiple climatic parameters or composite

 processes, such as atmospheric circulations (Boggs & Inoyue 2012; Boggs 2016; Forrest 2016; Halsch *et al.* 2021).

 To explain the climatic responsiveness of insects, several studies have shown the importance of a number of life history attributes (Robinet & Roques 2010; Halsch *et al.* 2021). In migratory insects, body size is predicted to be influential for climatic responsiveness as body metrics are key drivers of migratory performance in insects (Dingle et al 1980, Rankin & Burchsted 1992, Hu et al. 2018). The wintering stage has been shown to govern climatic responsiveness in lepidopterans. For example, a former study of ours has demonstrated that nocturnal moths (Noctuidae) hibernating as adults shift their time of emergence to earlier dates than species spending the winter as eggs, larvae or pupae (Végvári et al. 2015, Altermatt 2010).

 Apart from life history traits, a broad range of ecological and life history metrics of insects in general has been shown to be influential for climatic responsiveness, such as the number of generations per year, habitat association, overwintering strategy and responsiveness to specific climatic parameters (Halsch et al*.* 2021). Several investigations found that climate zone is a potential predictor of climatic responses in migratory insects, as the different climatic zones of the earth are differentially affected by climate change, inducing non-analogous processes in ecological networks (e.g. Woodward et al. 2010, Musolin & Saulich 2012).

 Although migratory insects exhibit immense taxonomic diversity and high variance in migration strategies, our knowledge of the environmental responsiveness of migratory strategies are extremely limited due to the lack of quantitative and long-term datasets.

 Thus, the aim of our study is to summarise the current state of knowledge on insect migration in relation to climatic processes. Our review is organised around the following research questions: (1) What are the levels of climatic complexity influencing behavioural responses of migratory insects? (2) Which life history traits influence climatic responsiveness?

**Methods**

#### *Literature search*

 We searched for research articles on insect migratory behaviour in connection with climatic changes. We used the Scopus [\(www.scopus.com\)](http://www.scopus.com/) database to generate a list of papers based on the following search terms: 'insect' and 'migration' and 'clima\*' including all papers published prior to June 2023, and expanded this list by searching for articles on known migratory species in Google Scholar.

 As a result, we found 446 papers matching the search terms. Using the abstracts, we narrowed down the list by excluding papers which only generally referred to relationships among migration and climate in insects, which resulted in a subset of 230 papers. Moreover, we also considered the migration type of the species during the selection of the papers for the analyses: we used only papers which considered seasonal migration (particular stages of annual journeys, following the recommendations of Dingle (2007)). Thus, dispersion and other types of movements among close populations were not considered as migration in this study. Finally, our search yielded 76 papers matching our criteria. In the next step, we divided these papers into two subsets: i) review articles providing additional general information on insect migration related to climate, which we excluded from further analyses as these lacked species-specific data and ii) research papers based on species- specific investigations on climatic responsiveness. We used the research papers to obtain data for further analyses.

### *Data analyses*

 First, we quantified the following metrics: number of papers from different years, number of investigated insect groups and geographic locations of the studies. Studies on the insects' responses to climatic processes were classified as those related to phenology or demography. As we found

 only three studies analysing demographical responses, we omitted these from further analyses, focusing only on the potential drivers of phenological response. In total, we used 57 papers in the further analyses (Appendix 1.) During paper search, selection and data retrieval, we included all species-specific records, irrespective of level of significance.

 In the next step, we created a database with the following response and proxy variables retrieved from papers. First, for each species we classified the climatic parameters which predicted a response, considering relationships between insect response and climatic predictors found in the original papers, independent of the level of significance (significant or not). Based on the array of dependent variables treated in the papers, in the next step, we classified the climatic responses into three main groups, using the species-level subsets of climatic parameters which relate to various levels of reactions to climatic complexity: response to i) the directness of the climatic parameter, classified as direct or indirect; ii) geographic scale of the climatic parameter, defined as local or continental and iii) complexity of the climatic response, defined as single, multiple, complex: "single" is assigned to species reacting to a single climatic parameter, "multiple" response indicates a reaction to several climatic parameters on local scales within the same study, referring to the same species, whereas "complex" indicates responsiveness to composite climatic patterns, such as atmospheric circulations, also within the same study (Appendix 2.) As potential predictors of these response variables, we considered i) body size (measured as wingspan in butterflies, provided as body length otherwise, provided in mm-s), which is widely accepted as a proxy of a large number of life history variables across the animal kingdom; ii) wintering stage was categorized as adult, egg, larva, or pupa; iii) climate zone describing the major climatic zones intersecting the distribution range, retrieved from taxon-specific literature and online databases (C FC Lt. Fraser 1936; Needham 2000; CABI 2022, GBIF.org 2022, Jonko 2022).

 As the study species are distributed over a broad taxonomic range of insects, we controlled for phylogenetic non-independence of the taxa by collecting order-level taxonomic information on the studied insects, as provided by the original papers.

### *Statistical analyses*

 We calculated the variance inflation factors (VIF) between all predictors to test for multicollinearity using the 'usdm' package in R. The VIF-values for all of the predictors were found to be < 1.30, thus considered as weakly correlated. As the next step, we fitted Generalised Linear Mixed Models (GLMM-s) on the i) climatic effect type (direct or indirect), ii) climate scale and iii) climate complexity as response variables, as a function of log-transformed body size, climate zone and wintering stage, treating study ID and insect order as random factors and considering binomial error family. We entered insect order as a random factor because this is recommended as a surrogate method for controlling for phylogenetic dependence without having a phylogenetic tree of finer resolution (Chamberlain et al. 2012).

 In the next step, we conducted a model selection procedure, considering each of the three models as full models separately, employing climatic effect, climatic scale and climatic complexity as response variables in the separate models. In the case of the weather complexity that has three levels, the 'glmer' package of the R statistical environment allows the use multi-level categorical response variables, thus performing multinomial regression; statistically, the three-level categorical response variable is treated as if it were a set of independent binary responses where each level of the response variable is considered a "success" with a certain probability, and the goal of the modelling is to model the probability of each level occurring as a function of the predictor variables 'lme4' package (Bates *et al.* 2015). After model fitting, the relative importance of life history traits (body size, wintering stage) and climate zone of distribution was calculated using information-theoretic model-comparison, considering all candidate models, which in our case meant the

 formulation of 8 candidate models for each response variable (Appendix 3.) Burnham and Anderson 2002). First, we obtained the values of Akaike's information criterion corrected for small sample size (AICc) and the corresponding Akaike weight of each model (ω). Second, calculated model- averaged parameter estimates (β), unconditional standard errors (SEu; Burnham and Anderson 2002), z-value of z-approximation and related p-values of each variable by the sums of their Akaike weights across all models. Akaike differences in the range 0–4 indicate substantial level of 202 empirical support of a given model, whereas  $\Delta i = 7$  implies very weak support (Burnham and Anderson 2002).

 GLMM-s were fitted applying the 'glmer' function available in the 'lme4' package (Bates *et al.* 2015) and model selection was carried out applying the 'MuMIn' package (Barton 2020); all of the packages are provided within the R Statistical Programming Environment (R Core Team 2022).

**Results**

*Primary database*

 Our dataset, compiled from 57 papers, consisted of records of 61 insect species classified into five insect orders, where each species occurred 1-5 times in the analysed investigations and the number 214 of species ranged between  $2 - 34$  in the insect orders (Fig 1, Appendix 2). The 57 original papers contained information for altogether 61 species, where several papers provided data for more than one species, which thus provided the 84 rows of the appendix table showing the original data (Appendix 2). The species in our study originated from five climate zones, where the number of 218 species ranged between  $10 - 17$  in the various climate zones (Fig 2., Appendix 2). The studied species originated from all continents with the exception of Antarctica and the number of species 220 per continent varied between  $1 - 25$  (Fig 3., Appendix 2). Considering the levels of significance of

 the climatic relationships reported in the papers, we found five studies that also reported non- significant results and only a single one reporting exclusively non-significant relationships; the subset of papers providing non-significant results referred to 10 species. However, the type of climatic response for the same metrics we applied for significant results (climatic effect, climatic scale and climatic complexity) was found to be constant; thus, due to the lack of variation for the studies with non-significant results, we could not include "level of significance" as a factor in the analysis.

### *Climatic drivers*

 Among the investigated climatic predictors, four key parameter groups emerged: temperature, precipitation and predictors describing the dynamics of the atmosphere, such as atmospheric pressure, wind and weather systems (Mak 2011). Out of these, temperature was demonstrated to be the most dominant climatic parameter, acting both directly and indirectly on climatic responsiveness of phenological patterns for the largest number of study species, distribution areas and trophic networks. For example, a study on the Northern Californian breeding population of the Monarch Butterfly reported a positive influence of warmer temperatures on the probability of Monarch presence in the breeding area (Lemoine 2015); however, the findings of that study indicate that host plant distributions predict Monarch distributions to a higher degree than climatic metrics. Similarly, Lemoine (2015) found that the ranges of both the Monarch and its food plant appear to be constrained by temperature, whereas the distribution of the host plant explains the majority of the variability found in Monarch observations. Another example of climatic influences mediated by host plant responsiveness to temperature changes is shown by aphids, for which the spring migration phenology has shifted in parallel to winter and early spring temperatures (Luquet *et al.* 2019). Direct temperature effects on population-level responses to climatic variability have been

 found in the noctuid moth *Autographa gamma*, the population sizes of which are closely related to winter temperatures of the overwintering areas (Chapman *et al.* 2012).

 Wind is an environmental driver that has increasingly been demonstrated to be highly influential for climatic responsiveness of migratory animals. For example, the migratory flights of the moth *Spodoptera frugiperda* are predominantly predicted by the speed and direction of wind (Westbrook *et al.* 2019). Similarly, shifts in wind direction and speed as a result of current climatic processes have been demonstrated to modulate migration routes of Desert Locusts as these insects are passive fliers (Cressman 2013).

 Considering correlations between migration strategy and climate, in 40 % of the papers we found direct relationships between the climatic variables and migration. For example, atmospheric circulations have been shown to play roles in the climatic responsiveness of *Nilaparvata lugens* and several *Spodoptera* species*,* whereas temperature was found to be the key predictor of climatic responses of *Autographa gamma* and several *Spodoptera* species (Westbrook *et al.* 2019, Torniainen & Mikonranta 2018, Hu et al. 2019). In the other 60 % of the selected papers, we detected only indirect connections, which were mainly linked to food plant phenology in response to climatic changes. During the detailed analysis we uncovered that the indirect effect of climatic 261 changes was detected mainly  $(70\% \text{ of N=18 studies})$  in lepidopterans.

## *Associations among climatic responsiveness, life history and ecological predictors*

 As the model selection procedure performed on the full generalised linear mixed models requires a dataset without missing values, the omission of records where at least one ecological or life history predictor was lacking – which happened in several cases – the sample size correspond to the number of records of the filtered dataset, which is 72 in our analyses. The directness of the climatic effect was predicted by climate zone(s) and marginally significantly by body size: species distributed in regions with temperate as well as tropical climates were largely indirectly influenced

 by climate parameters, whereas insects with distribution areas in colder regions responded more 271 strongly to direct effects (model-averaged GLMM,  $N = 72$ ,  $b = -27.5318$ ,  $p = 0.045$ ; Table 1, Fig 4); larger-bodied insects tended to respond more to indirect climatic effects, whereas smaller species 273 were more responsive to direct climatic effects (model-averaged GLMM,  $b = 14.0235$ ,  $p = 0.0389$ ; Table 1, Fig 5, Appendix 3).

 Climatic complexity was strongly associated by body size: with decreasing body size, insects were increasingly responsive to multiple climatic patterns, reacting increasingly strongly to simple and 277 complex patterns (model-averaged GLMM,  $N = 72$ ,  $b = 8.7468$  p = 0.0003; Table 2, Appendix 3).

 Climatic scale was unrelated to any of the model predictors (body size, wintering stage and climate 279 zone; model-averaged GLMM,  $N = 72$ ,  $p \ge 0.9478$  for all cases; Table 3, Appendix 3).

 The phylogenetic signal in the global model measured as the variance of the insect order treated as 281 random term was found to be low:  $S^2 \pm SD = 8.58 \pm 2.929$ .

**Discussion**

 Our key findings indicate that migratory insects respond to various levels of complexity in climatic patterns and these responses are predicted by life history and ecological traits: i) responses to climatic effect type were predicted by climate zone(s) of the distribution area, whereas ii) response to climatic complexity was predicted by body size. We discuss these relationships in detail.

 Our result that species distributed in regions with colder climates responded more to direct effects than insects with distribution areas in warmer regions can be explained by the following, mutually non-exclusive hypotheses. First, climatic processes are less complex in the cold and/or temperate and arctic zones than in the temperate and/or tropical regions (Lydolph & Temple 1985), which is thus predicted to select for responsiveness to direct climatic effects in insects. Alternatively, the annual activity period of insects distributed in colder regions is substantially shorter and constrained  to the warmer months of the year than those occupying hotter climatic zones which induces the emergence of more direct responses to weather parameters such as temperature changes. For example, spring emergence in many species are driven by direct climatic cues (Boggs 2016).

 Our finding that larger-bodied insects were more responsive to multiple local climatic patterns, whereas smaller-bodied species reacted more to both simple and complex local patterns invokes two mutually non-exclusive hypotheses. First, the decreased ability of smaller-bodied insects to perform active migratory movements in the air as compared to larger insect predicts that species with smaller body size and migratory ranges are more responsive to simple climatic parameters than to complex ones. Alternatively, smaller species migrating on continental scales may benefit from evolving navigation senses using complex environmental cues leading to optimised navigation decisions (Aralimarad *et al.* 2011). Indeed, larger insects perform better in active flight; one of the hypotheses explaining this pattern states that larger insects are less dependent upon evolving navigational mechanisms supporting passive flight mechanisms, which might be highly important for the survival of smaller insect species during migration (Menz *et al.* 2022). However, to test these hypotheses, we need to have more data on the migration distances of the studied species.

 The number of research papers analysing relationships among insect migration and climatic processes has been growing in the past few decades, deepening our knowledge on the environmental background of migratory mechanisms in ectotherms. This is facilitated by the fast accumulation of information on climatic changes during this period. We found that the majority of the papers in consideration confirm that migratory behaviour in insect species is impacted by climatic processes, however, the causal relationships among the underlying mechanisms are still far from understood. Further, a significant component of the complexity of determining how changes in climate may impact migratory species is driven by the identification of the potential climate variables which are most relevant to the biology of the given species. Indeed, insects are sensitive to particular meteorological conditions to various degrees and this sensitivity may vary over time

 (including various aspects of their life cycle) and across the range of the species (Chambers 2009). Furthermore, interspecific differences can also be detected in sensitivity to environmental factors such as temperature, therefore warming can increase or decrease synchronization between insects and their food plants and/or natural predators (Forrest 2016).

 Considering the investigated climatic predictors, temperature was the most commonly studied climatic parameter, which affects migration in different ways. For example, Bradshaw & Holzapfel (2010a, b) investigated the geographical and seasonal patterns in light and temperature in the temperate zone, and how climate change influences migration patterns and the actual and potential biotic responses by insects to climate warming. They found that across a large taxonomic range, each individual shows genetically driven responses to temperature, which induce irrevocable physiological decisions whether to initiate migration or not. Future environmental conditions then determine the fitness consequences of that decision. Specifically, physiological processes drive functional relationships between the genotype and the phenotype that ultimately confronts the environment and influences individual fitness. As wild animals face environmental variability and cope with that variation through physiological adjustments at the behavioural, hormonal, cellular, and biochemical levels.

 Considering response variables, we found that phenological shifts were the most commonly found reaction to climatic changes. Related studies indicate that changes in phenology can influence species interactions (including pollination), population viability, water cycles, carbon sequestration, human health (allergenic disorders, vector borne diseases, pest insect control), tourism and recreation, transport (e.g. bird migration and aircraft collisions), timing of management activities and productivity in agriculture (Chambers 2009). Given the often strong relationships between changes in climate and changes in phenology, phenological measures can also be used as proxies for climate change. Most of the knowledge about changes in phenology comes from the Northern

 Hemisphere mainly from Europe alone (Chambers 2009) limiting our understanding of the diversity of organismal response to climate change.

 Numerous effects of climate change on spring phenology of plants and animals have been documented (Wolkovich *et al.* 2012; Parmesan *et al.* 2013). We also noticed that in investigations analysed in our study, the changes in insect phenology were always monitored in spring. Interestingly, autumn seemed to be a neglected season in climatic studies conducted in temperate and arctic ecosystems. A recent study analysing this pattern (Gallinat *et al.* 2015) showed that leaf senescence and bird migration are the most frequently examined responses in autumn, and insect diapause and fruit ripening have so far received only moderate interest.

 Our study might be biased by the latitudinal distribution of study locations, as the majority of the investigations have been conducted in the Northern Hemisphere, which tendency is also found in avian studies of climatic responses (Dunn & Moller 2019). This calls for a more active sampling at southern latitudes which might be important for outlining management plans for endangered species. Additionally, the publication bias towards publishing mostly significant results – also found in our study – might affect the results of meta-analyses, which has also been reported in ecological studies (Lortie et al. 2007). To overcome this problem, we encourage ecologists to publish non- significant results, which is expected to deepen our understanding of fundamental ecological issues. We also detected several knowledge gaps in climatic investigations of migratory insects, which when addressed would substantially contribute to clarifying some causal mechanisms. Out of these, we emphasise the small number of studies i) analysing relationships among physiological adaptation to migration and climate and ii) the genetic and epigenetic background of climatic fingerprints in migratory behaviour.

 Finally, we recommend to investigate the so far underestimated effects of changes in a wider range of potential environmental predictors such as climatic teleconnection networks – defined as large-scale atmospheric and oceanic interactions that link different regions of the Earth's climate system,

 which involve the exchange of energy, moisture, and momentum between distant locations, influencing weather patterns and climate variability on a global scale – solar radiation, humidity, and levels of greenhouse gases, which are expected to modulate climatic responsiveness in migratory insects (Robinet & Roques 2010). Climatic extremities also need more focus in climatic ecology, as the increasing frequency of arid periods, storms, extreme cold events in spring and heat shock days have already been shown (Forrest 2016) to exert strong effects on insect migration, by blocking or altering migration routes.

 To conclude, we show that migratory insects respond to various levels of complexity in climatic processes, and this responsiveness is i) predicted by life history and ecological proxies; and ii) governed by a substantially wider array of environmental predictors than demonstrated in vertebrates.

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## 564 **Tables**

565 Table 1. Model selection table for the GLMM fitted on the climatic effect as response variable. 567 Significant relationships are marked in bold.  $β$  indicates parameter estimates, SEu stands for unconditional SE-s. unconditional SE-s.

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573 Table 2. Model selection table for the GLMM fitted on the climatic complexity as response 575 variable. Significant relationships are marked in bold. β indicates parameter estimates, SEu stands 576 for unconditional SE-s. for unconditional SE-s.

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**586 Table 3**. Model selection table for the GLMM fitted on the climatic scale as response variable. β indicates parameter estimates, SEu stands for unconditional SE-s. indicates parameter estimates, SEu stands for unconditional SE-s.

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# 600 **Figure legends**

601 Fig. 1. a) Taxonomic distribution of migratory insects, indicating the number of species classified 603 into insect orders; **b)** Climatic zonal distribution of migratory insects, indicating the number of species occurring in major climatic zones; **c**)Spatial distribution of migratory insects, indicating the 605 number of species occupying various continents.

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608 Fig 2. Relationship between climatic effect (direct or indirect) and climate zone (defined as the major climatic zones intersecting the distribution range) for migratory insects.

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611 Fig 3.Relationship between climatic effect and log-transformed body size for migratory insects, showing the level of significance for the GLMM fitted on the climatic effect as a function of bo

showing the level of significance for the GLMM fitted on the climatic effect as a function of body 613 size.



 Fig 2. 



 

