

Migration in butterflies: a global overview

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ABSTRACT

Insect populations including butterflies are declining worldwide, and they are becoming an urgent conservation priority in many regions. Understanding which butterfly species migrate is critical to planning for their conservation, because management actions for migrants need to be coordinated across time and space. Yet, while migration appears to be widespread among butterflies, its prevalence, as well as its taxonomic and geographic distribution are poorly understood. The study of insect migration is hampered by their small size and the difficulty of tracking individuals over long distances. Here we review the literature on

migration in butterflies, one of the best-known insect groups. We find that nearly 600 butterfly species show evidence of migratory movements. Indeed, the rate of ‘discovery’ of migratory movements in butterflies suggests that many more species might in fact be migratory. Butterfly migration occurs across all families, in tropical as well as temperate taxa; Nymphalidae has more migratory species than any other family (275 species), and Pieridae has the highest proportion of migrants (13%; 133 species). Some 13 lines of evidence have been used to ascribe migration status in the literature, but only a single line of evidence is available for 92% of the migratory species identified, with four or more lines of evidence available for only 10 species – all from the Pieridae and Nymphalidae. Migratory butterflies occur worldwide, although the geographic distribution of migration in butterflies is poorly resolved, with most data so far coming from Europe, USA, and Australia. Migration is much more widespread in butterflies than previously realised – extending far beyond the well-known examples of the monarch *Danaus plexippus* and the painted lady *Vanessa cardui* – and actions to conserve butterflies and insects in general must account for the spatial dependencies introduced by migratory movements.

Key words: butterfly, butterfly migration ecology, conservation, insect movement, Lepidoptera, migratory butterflies, monarch, painted lady, seasonal movement.

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I. INTRODUCTION

Migration is a widespread phenomenon among animals, particularly in vertebrates, insects, and marine invertebrates (Alerstam, Hedenström & Åkesson, 2003; Dingle, 2014). Animals that migrate range in size from tiny insects to blue whales, and comprise a remarkable proportion of animals worldwide, with that fraction constantly increasing as we learn more about animal movements (Cioc, 2009; Dingle, 2014). Distances travelled vary greatly, with some species making just local movements (Denno & Grissell, 1979; Denno *et al.*, 1980; Denno, Douglas & Jacobs, 1985), and others traveling thousands of kilometres between continents and even hemispheres (Alerstam *et al.*, 2003; Chapman, Reynolds & Wilson, 2015). From a conservation viewpoint, the often–vast geographic range sizes of migratory species might be assumed to protect them from extinction risk, yet migratory species can be acutely prone to threats occurring at key sites where they congregate at some point in the annual cycle (Koh & Sodhi, 2004; Wilcove & Wikelski, 2008; Flockhart *et al.*, 2015; Runge, 2015; Runge *et al.*, 2015; Stenoien *et al.*, 2016; Dhanjal–Adams *et al.*, 2017; Chowdhury *et al.*, 2020; Gao *et al.*, 2020). For example, habitat destruction and fragmentation, coupled with intense human predation led to the extinction of at least three formerly widespread and common migratory species: the Rocky mountain grasshopper *Melanoplus spretus*, the

passenger pigeon *Ectopistes migratorius*, and Eskimo curlew *Numenius borealis* (Brower & Malcolm, 1991; Bucher, 1992; Hopkins, 2005; Harris *et al.*, 2012). Migratory connectivity between habitat patches may complicate the planning of conservation management activities, because it is necessary to account for spatial dependencies among sites when evaluating the benefits of alternative conservation actions (Brower & Malcolm, 1991; Martin *et al.*, 2007; Miller *et al.*, 2012; Iwamura, Fuller & Possingham, 2014; Flockhart *et al.*, 2015; Dhanjal-Adams *et al.*, 2017).

Migration enables individuals to find and occupy more favourable habitats as or before the current habitat deteriorates, often (although not always) on a regular temporal schedule.

Individuals exhibit morphological, physiological, and behavioural traits that enable migration to occur (Berthold, 1999; Åkesson & Hedenström, 2007). For example, many migratory animals have an internal ‘biological clock’ (Gwinner, 1996) that schedules the various components of the migration (Liedvogel, Åkesson & Bensch, 2011). To find their species- or population-specific seasonal quarters, migrating animals need to combine elements of temporal and spatial information into a spatiotemporal migration programme (Åkesson & Hedenström, 2007).

Insect migration is much less well studied than migration in birds and other vertebrates, despite its potentially huge importance to ecological resource flows and ecosystem services globally (Hu *et al.*, 2016; Menz *et al.*, 2019; Wotton *et al.*, 2019; Satterfield *et al.*, 2020).

Migration occurs in all major insect taxonomic groups, although its prevalence varies markedly among species, even within closely related clades (Drake & Farrow, 1988; Dingle, 2001; Dingle *et al.*, 2005). Migratory species occur in all functional groups of insects: pollinators, herbivores, predators in the broad sense, decomposers, and of course, pests – insects that are vectors of human, livestock and plant diseases, as well as those that directly

damage crops (Walther *et al.*, 2002; Dingle, 2006, 2009; Reynolds, Chapman & Harrington, 2006). Migration is the key process in the ecology and population dynamics of many insect species, yet perhaps as a consequence of their small size and general lack of mass aggregation in comparison with vertebrates, the migration of insects remains mostly unnoticed and, with only a few exceptions, poorly studied (Chapman *et al.*, 2015; Gao *et al.*, 2020; Satterfield *et al.*, 2020). Yet new technologies and the advent of mass-participation citizen science are beginning to document the large-scale movements of insects in ways that have hitherto been unavailable (Stefanescu *et al.*, 2013; Cohen *et al.*, 2019; Knight *et al.*, 2019; Juhász *et al.*, 2020). Here we review and synthesise what is known about migration in butterflies, and outline some of the key information gaps.

Butterflies are one of the better-studied groups of insects; generally, there is a good understanding of their ecology, and we know that the degree of connectivity between fragmented populations is critical to effective conservation (Brower & Malcolm, 1991; Hanski & Thomas, 1994; Baguette, Petit & Quéva, 2000; Hanski, Alho & Moilanen, 2000; Koh & Sodhi, 2004; Gao *et al.*, 2020; Chowdhury *et al.*, 2021). About 19% of all bird species migrate (Kirby *et al.*, 2008; Somveille *et al.*, 2013), but perhaps surprisingly, the number of migratory butterfly species, often dubbed the ‘birds of the insect kingdom’, remains unknown. There are reports of migration across various butterfly families, suggesting that at least a few hundred species might migrate, but given the difficulty of studying the phenomenon it is plausible this is vastly underestimated. Williams (1930) summarised all existing research at the time of writing and reported 217 migratory butterflies. This work is now 90 years old, and there is no recent comprehensive review of the prevalence of butterfly migration.

Here we conduct such a review, collating and summarising published descriptions of butterfly migrations globally, highlighting key aspects of the migration phenomenon in butterflies, analysing previous findings, and pointing out major knowledge gaps. We ascertain, to the extent that current knowledge allows, and based on multiple lines of evidence, which butterfly species are migratory, provide lists by zoogeographic region, show where and how butterfly migration has been studied, and evaluate potential drivers of migration. We hope this work will stimulate greater scrutiny and assessment of migratory movements made by butterflies, with a view to improving ecological understanding and planning systematically for butterfly conservation.

II. BUTTERFLY MOVEMENTS

Butterflies are relatively large-bodied for insects, reasonably well described taxonomically, often colourful, and usually diurnal (compared with the much larger number of moth species), and they are more likely to be observed and recorded than any other insect group (New, 1997; Hickling *et al.*, 2006). Butterflies are known to exhibit the full range of movement types, from extreme sedentariness (Ehrlich, 1986; Walther *et al.*, 2002) to some of the longest-distance insect migrations, in which individuals cover thousands of kilometres or perform complex intergenerational relays over similar distances (Stefanescu, 2001; Mouritsen *et al.*, 2013b; Stefanescu *et al.*, 2013; Runge *et al.*, 2014). Some *Euphydryas* checkerspot butterflies are extremely sedentary, remaining restricted to localised colonies and with only short-distance dispersal (Ehrlich & Gilbert, 1973), although movement patterns vary markedly among species (Brown & Ehrlich, 1980), and among populations of the same species (Gilbert & Singer, 1975). Edith's checkerspot *Euphydryas editha* is the least vagile checkerspot butterfly in the western USA, although it too disperses when disturbed or when

habitat conditions become suboptimal (Ehrlich, 1986). By contrast, monarchs *Danaus plexippus* travel about 5000–6000 km to complete their annual migration over 3–5 generations (Brower, 1995), and regularly move between patches of larval food plant during the breeding season (Grant *et al.*, 2018), although they frequently become ‘sedentary’ after colonising small islands (Freedman *et al.*, 2018). Painted ladies *Vanessa cardui* migrate more than twice as far (up to about 15,000 km) in round-trip migrations from Northern Europe to West Africa over at least 6–7 generations (Pollard *et al.*, 1998; Stefanescu *et al.*, 2013; Talavera *et al.*, 2018).

Butterfly movements fall into two main types: (1) flying within, or (2) going beyond their local habitat or some other circumscribed area (equivalent to the home range of vertebrates). Butterflies showing the latter type of behaviour are often considered migrants (Dingle, 1996, 2009, 2014; Nesbit *et al.*, 2009). The major difference between migrant and non-migrant individuals is that the former generally do not get distracted by appetitive cues such as food items and particularly sexual signals (Dingle, 1978; Chapman & Drake, 2016), although migratory butterflies often pause their journey to collect nectar and then resume flight towards their destination (Dingle, 2009). In vertebrates, migration is usually taken to mean an outward and return journey by the same individual (Dingle & Drake, 2007), yet due to their short lifespan, the pattern in butterflies is for successive generations to execute long-distance directional movement resulting in an overall change in the distribution of a population in response to an environmental cue (Baker, 1969; Alerstam *et al.*, 2003; Milner-Gulland, Fryxell & Sinclair, 2011).

The definition of migration varies from author to author, but a common thread among most definitions is that individual behaviour can be used to differentiate migration from other movement behaviours (Dingle & Drake, 2007). Kennedy (1985, p. 8) proposed a widely

used definition based upon behavioural aspects: “Migratory behaviour is persistent and straightened-out movement effected by the animal’s own locomotory exertions or by its active embarkation on a vehicle. It depends on some temporary inhibition of station-keeping responses but promotes their eventual disinhibition and recurrence”. Not all migrants will show the same suite of characteristics, and flexibility around characterising and diagnosing the phenomenon is needed (Kennedy, 1985).

In comparison with many vertebrates, migrations in butterflies may seem rather atypical, which has led to a diversity of viewpoints on what constitutes migration in butterflies. In this review, we adopt the broad behavioural definition quoted above, recognising that it can only be applied with certainty to a small number of species. However, we argue that this reflects the lack of research (and the resultant lack of data and knowledge) on these butterflies, rather than a failure of the definition itself. As pointed out by Gatehouse (1997), there remains a lack of research on the behavioural and physiological aspects of migration with respect to Kennedy’s (1985) definition. We hope that our review will stimulate such research to determine whether particular species or individuals are in fact migrants. Some movement ecology researchers, particularly those with experience of ‘classic’ vertebrate migration systems might consider some of the butterflies we classify as migratory as better classified as dispersive species; however, we adopt this broader view to establish a first framework for future studies of butterfly migration.

Partial migration, whereby part of any population migrates while the rest remains resident, (Chapman *et al.*, 2011a; Malcolm & Slager, 2015; Slager & Malcolm, 2015; Ruiz Vargas *et al.*, 2018; Menz *et al.*, 2019) is likely to be common in butterflies. For example, Vander Zanden *et al.* (2018) collected 33 monarchs in south Florida; isotopic analysis showed that 48% of the individuals originated from further north (i.e. they were recent immigrants), while

52% had developed locally (i.e. they were residents). As butterflies are generally short-lived with rapid generation times, the migratory cohorts are more likely to be separated temporally rather than spatially (as is typical of vertebrate partial migrants; Chapman *et al.*, 2011b), and such situations have been termed ‘sequential partial migration’ (Ruiz Vargas *et al.*, 2018). This category of partial migration may well prove to be the most common type of migration exhibited by butterflies. The key to resolving such issues will be to approach the study of migration from more than just observations of numbers flying past a fixed location, the main approach used to categorise migration in most species to date (see Section IV).

III. LITERATURE SEARCH

We searched *Google Scholar*, *PubMed*, and *Web of Science* to determine which butterfly species have been considered migratory according to at least one or more lines of evidence. Initially, we used ‘butterfly migration’ and ‘migratory butterflies’ as search terms in *Google Scholar* and carefully scanned the first 1000 search results for relevant papers. We then searched *Google Scholar* for specific information on the migratory status of each species (*X*) identified during the initial search using the search term ‘*X* migration’. In *PubMed*, the search terms ‘butterfly migration’ and ‘migratory butterflies’ yielded a total of 228 journal articles. A *Web of Science* search using ‘TS=(Butterf* AND migr*)’, yielded 1,297 publications. In both cases we followed up each suspected migratory species with a specific search. We also checked the bibliography of each article to identify other relevant literature. Many authors publish on butterfly migration in local journals or newsletters (e.g. *Atalanta*, *Phygea*, *Bulletin of the Allyn Museum*, and *Journal of the Bombay Natural History Society*), yet during our literature search using the above search engines, we rarely located relevant work in these journals. Therefore, we carefully checked all issues of these publications to locate butterfly

migration-related publications. Eventually, during our literature search in both Google Scholar and Web of Science, we found repetition of the same literature and started to come across hundreds of non-relevant works, suggesting we were approaching saturation, at least of the readily available literature. Together, from all sources we identified 422 articles, 36 books, 9 book chapters, 2 book reviews, 10 letters, 3 meeting abstracts, 16 notes, 43 review articles, 23 web articles, 5 theses and 9 proceedings (marked with asterisk (*) in the reference section). Our searches were conducted up to December 2019 and we primarily considered articles written in English; however, if we identified relevant publications in other languages, these were translated.

IV. EVIDENCE FOR MIGRATORY STATUS

We focused on the primary literature on observations of butterflies migrating or for specific instances in which migratory behaviours were inferred, and built a list of methods that have been used to ascribe migratory status (Table 1). These fell into the three major groups of experimental, physiological and population evidence. General assertions that a species was migratory were not sufficient to consider that migration had been proved for a particular species. Instances of directional flights in which individuals appeared undistracted strongly suggest that the butterflies were migrating, and were included (Dingle, 2001; Dingle & Drake, 2007). In other instances where there was no mention of individual behavioural characteristics, migration could still be inferred if there was documentation of butterflies occurring in the absence of larval food plants, performing nomadic or seasonal movements, where large numbers were recorded roosting or imbibing nectar, and where there was evidence of formation of overwintering aggregations at specific sites that are strongly suggestive or indicative of migration (see Table S1 for full details).

Table 1. Methods that have been used to attribute migratory behaviour to butterflies with example references for each. See Table S1 for full details of species studied using each of these methods.

| Evidence (broader categories) | Evidence (specified) | Description | References |
|--------------------------------------|---|---|---|
| Physiological | Larvae are incapable of developing due to unfavourable seasonal climate | The environment is not suitable for the larvae to complete development | Clements (2012) |
| Physiological | Extended pre-oviposition period | Female butterflies delayed egg maturation, suggesting time available to migrate rather than reproduce | Johnson (1963) |
| Physiological | Larval food plant absent | Larval food plants were absent where the butterfly was seen, suggesting they were flying through and not breeding | Braby (2005) |
| Physiological | Breeding population not established | Butterfly recorded in an area where there was no breeding record | Braby (2000) |
| Experimental | Mark–release–recapture | Marked individuals displaced away from mark site and recaptured along ‘flight path’; marked individuals recaptured a long way from marking site along flight path | Urquhart (1960); Urquhart & Urquhart (1976); Matter <i>et al.</i> (2004) |
| Experimental | Trap catches | A trap built along the migratory route was used to catch migrant butterflies and identify their migratory direction | Walker (1985 <i>b</i>) |
| Experimental | Tethered flight measurements | Used to establish orientation and potential displacement under various conditions | Mouritsen & Frost, (2002); Reppert <i>et al.</i> (2004); Mouritsen (2018) |
| Experimental | Amplified fragment length polymorphism (AFLP) | The occurrence of distinct genetic types can be used to understand whether long-distance movements occur, and to link breeding and non-breeding grounds. | Brattström <i>et al.</i> (2010 <i>a, b</i>) |
| Experimental | Radar observations | Provide direct records of insects flying at high altitude | Drake & Farrow (1985); Chapman <i>et al.</i> (2011 <i>b</i>) |
| Experimental | Flight in beneficial compass direction | Establish vanishing angles, direction of movement, orientation, etc. | Srygley (2001); Srygley & Oliveira (2001); Srygley <i>et al.</i> (2006) |
| Experimental | Stable isotopes | Stable isotope (dD) used to determine probable geographic origins of migrants | Wassenaar & Hobson, (1998); Brattström <i>et al.</i> (2010 <i>a, b</i>). |

| Evidence (broader categories) | Evidence (specified) | Description | References |
|--------------------------------------|--|--|---|
| Experimental | Maintaining constant compass course by using wind Drift Compensation | Flights were compensated with respect to wind speed/direction | Srygley (2001); Srygley & Dudley (2008) |
| Population evidence | Mass movement | Many butterflies moving in the same direction at the same time, and that were apparently undistracted by appetitive cues | Shapiro (1982, 1986); Stefanescu (2001); Burwell <i>et al.</i> (2011) |

We identified 13 lines of evidence that have been used to diagnose butterfly migration and grouped them into three broader categories: experimental (8 lines of evidence), physiological (4), and population evidence (1) (Table 1). For each species, we collated sources that provided evidence of migration status, categorised into one of these 13 types (Table 1). The number of methods to assess migratory status of a butterfly varied markedly (Table S1). We found only a single line of evidence for 92% of species classified as migratory (521 of the 568), two or three lines of evidence for 36 species, and four or more lines of evidence for only 10 species (all in Pieridae and Nymphalidae). We found a further 23 species where authors asserted that a species was migratory but did not provide any evidence; we do not consider that migration has been proved in these species and exclude them from this review. It is clear that most work on butterfly migration has focused primarily on a small suite of species, with only 46 species having more than one line of evidence for migration in the literature. There remains a great deal of work to do on the migration of most butterfly species. Although direct experimental work is the best indicator of migratory behaviour, most authors used changes in distribution (population evidence), or records of behaviour (Fig. 1) to infer migration. For example, many authors consider a butterfly to be a migrant if mass movement

has been noted; this line of evidence accounts for the diagnosis of migration in more than 90% of cases. Analysis of amplified fragment length polymorphism (AFLP) or similar approaches has been used to determine migratory status only for a single species (red admiral *Vanessa atalanta*) despite the fact that this method is commonly used to determine population structure and migration among bird species (Bensch, Åkesson & Irwin, 2002; Bensch & Åkesson, 2005; Brattström, Åkesson & Bensch, 2010a; Brattström *et al.*, 2010b). Flight chamber experiments involving tethered flight have been used for only five species (see Table S1).

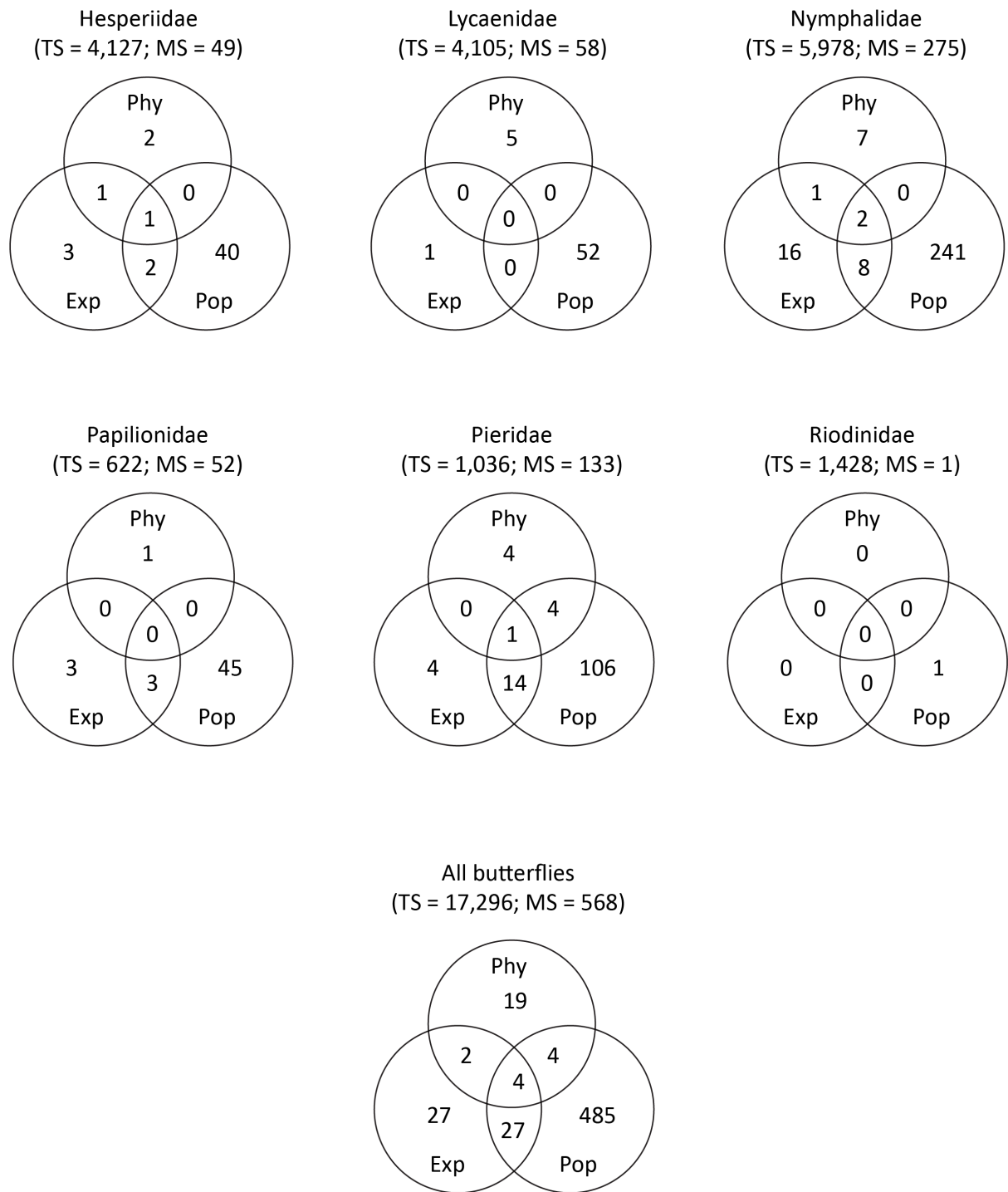


Fig. 1. Methods used to ascribe migratory status to butterflies. Here, TS = Total identified species, MS = Total recorded migratory species, Phy = physiological evidence; Exp = experimental evidence; Pop = population evidence. The species number beside each family is the total number of species in the world and numbers in the circles indicate how many

species have been diagnosed as migratory with that particular method or combination of methods. The total number of butterflies in each subfamily is derived from an online data repository (<https://www.learnaboutbutterflies.com/World%20Census.htm>). Note that many species have been diagnosed as migratory based on only a single category of evidence.

V. PREVALENCE OF MIGRATION

In total, we found evidence for migration in 568 species of butterflies belonging to 221 genera, from 21 subfamilies, across all six major families (see Table S1). At the family level, Pieridae had the greatest proportion of migrants, with 13% (133 species) being documented as migratory, followed by Papilionidae (8%; 52 species), Nymphalidae (5%; 275 species), Lycaenidae (1%; 58 species), Hesperidae (1%; 49 species), and finally only one of the 1428 species in the Riodinidae (Fig. 1). At the sub-family level, Libytheinae (Nymphalidae) contains the highest proportion of butterfly migrants (nine species, 53%), while Pierinae (Pieridae) contains the highest number of migrants (80 species, 11%). Interestingly, migrants appear more prevalent in smaller subfamilies (Fig. 2), but it is unclear whether this represents differences in the extent to which the different subfamilies have been studied or genuine differences in the biological prevalence of migration in some phylogenetic clades.

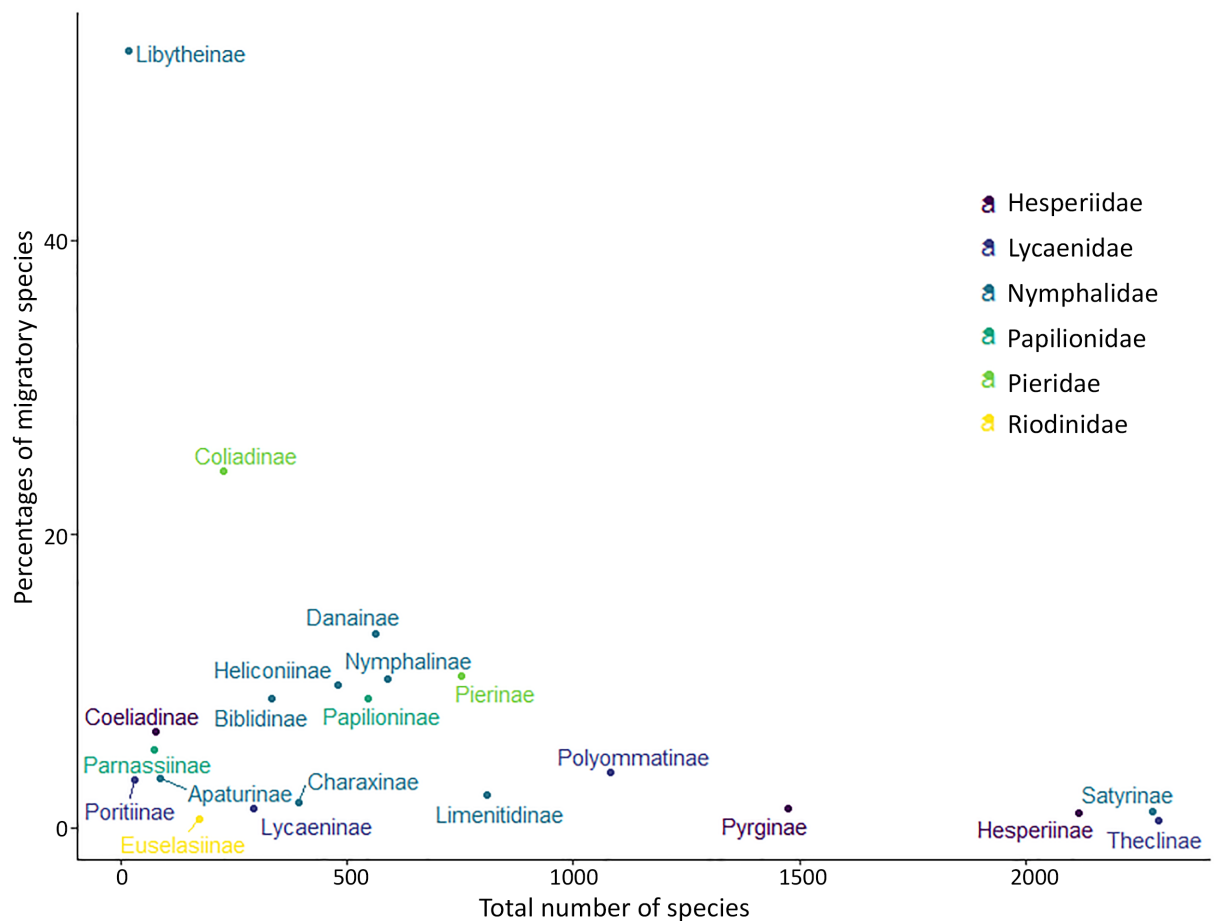


Fig. 2. Prevalence of migratory species among butterfly subfamilies. The total number of butterflies in each subfamily is from <https://www.learnaboutbutterflies.com/>.

It is clear that butterfly migration is a widespread global phenomenon, although the number of recorded migrants varies markedly from continent to continent and from region to region (Fig. 3). While only 3% of described butterfly species globally have been diagnosed as migratory to date (Fig. 1), the proportion varies markedly from one zoogeographic region to another. Migratory butterflies are mostly recorded from the Holarctic (Nearctic and Palearctic) region (12% in each); while in the Neotropical and Afrotropical regions together, around 5% of the butterflies are migratory, 7% from the Australian region, and roughly 8%

are migratory in the Oriental region (Fig. 3). These results are likely to reflect substantial regional bias in research attention and many migrants might remain undetected.

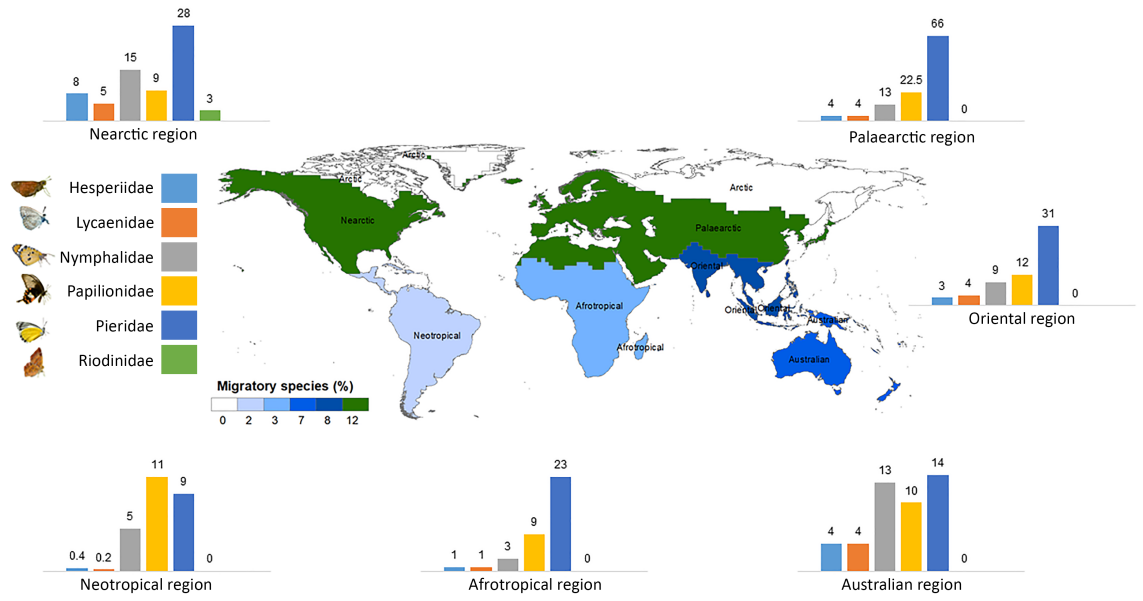


Fig. 3. Prevalence of butterfly migration across the world. Colours on the map indicate the percentage of all described butterflies for each zoogeographic region that are considered migrants. The bar charts indicate the percentage of described butterflies in each zoogeographic region that are considered migrants, separated according to family.

Migratory butterflies were recorded from all continents except Antarctica (which has no butterflies). Most butterfly migration was recorded from predominantly tropical countries (e.g. Australia, India, and Sri Lanka had the highest total numbers of migratory species), indicating that this phenomenon is not restricted to strongly seasonal high latitudes. However, when expressed relative to the total butterfly species present, higher latitude temperate or cooler countries have higher proportions of migratory species: 12% of Palaearctic butterflies are migratory compared to 3% of the butterflies from the Afrotropical region (Fig. 3). In an analysis of the distribution of Australian migratory and non-migratory butterflies, Dingle *et al.* (2000) found migrants to be more common in arid areas.

The geographic spread of species that migrate varied among families. In the Papilionidae, migratory species were mostly documented in the Palaearctic (23%), while for the Nymphalidae migrants were most common in the Nearctic (15%). In all five regions globally, migration was most prevalent in the Pieridae, while only one species in the Riodinidae was reported as a migrant (from the Palaearctic) (Fig. 3). Although 13% of the world's pierids are migratory, the proportions are substantially higher in some regions (66% of Palaearctic pierids, 31% of Oriental pierids, and 28% of Nearctic pierids are migratory) and negligible in others (Fig. 3).

There was a clear disparity between documentation of migration, and the global distributions of species in which migration has been documented. This presumably indicates either a

spatial bias in where migration is studied, or that these species do not migrate in all parts of their geographic distribution. Records from Afrotropical and Neotropical regions were scarce. In addition, we could only find evidence for a single migratory species of Riodinidae (*Euselasia euboea*; Williams, 1930). Riodinid butterflies are generally restricted to specific microhabitats and may be spatially rare with low population densities, even where they have wide distributions (DeVries, Chacon & Murray, 1992; Campbell & Pierce, 2003; Siewert *et al.*, 2014; Espeland *et al.*, 2015). It would be interesting to further investigate the migration of Riodinidae since we found only one reference (*Euselasia euboea*; Williams, 1930) documenting an instance of migration in this family. Future studies could investigate whether there are particular ecological or genetic characteristics that constrain riodinids in terms of migration.

Migration has been observed across the geographic range of several widespread species, with up to 30% of butterflies recorded as migrants from more than one region. *Vanessa cardui* was recorded as a migrant in six regions; migration of *Hypolimnas misippus*, *Ascia monuste*, and *Pieris rapae* has been recorded from five regions; twelve species (*Belenois aurota*, *Colias croceus*, *Colias hyale*, *Danaus chrysippus*, *Danaus plexippus*, *Hypolimnas bolina*, *Junonia orithya*, *Lampides boeticus*, *Libytheana carinenta*, *Nymphalis antiopa*, *Pieris brassicae*, and *Vanessa atalanta*) migrate in four regions.

For 24 species, we found inconsistent records: one author specifically noting sedentariness in contrast to another author considering the same species to be a migrant elsewhere (Table 2). For example, both *Heliconius erato* (Turner, 1971) and *H. ethilla* (Ehrlich & Gilbert, 1973) have been described as sedentary, although *H. ethilla* and other *Heliconius* species have been recorded elsewhere as taking part in large-scale migrations (Williams, 1930; Beebe, 1949a, b; Beebe, 1950a,b).

Table 2. List of species that have been recorded as both migratory and non-migratory/resident.

| Species | Recorded as migratory | Recorded as non-migratory |
|------------------------------|---|----------------------------------|
| <i>Acraea terpsicore</i> | Williams (1927, 1930); Larsen (1988); Dunn & Petrie (2017) | Bhaumik & Kunte (2018) |
| <i>Aporia crataegi</i> | Williams (1930); Baker (1969); Larsen (1975); Beshkov (1996a); Bolotov <i>et al.</i> (2013) | |
| <i>Araschnia levana</i> | Bolotov <i>et al.</i> (2013) | Parmesan <i>et al.</i> (1999) |
| <i>Argynnis adippe</i> | Baker (1969) | Parmesan <i>et al.</i> (1999) |
| <i>Celastrina argiolus</i> | Williams (1930); Williams <i>et al.</i> (1942); Baker (1969); Cook <i>et al.</i> (2001) | Parmesan <i>et al.</i> (1999) |
| <i>Colias eurytheme</i> | Shapiro (1973, 1980a); David (1982); Ferris (1989, 1993); Fleishman <i>et al.</i> (1999) | Ferris (1988) |
| <i>Colias philodice</i> | Fenton (1919); Williams (1930); Clark (1931) | Ferris (1988) |
| <i>Colotis fausta</i> | Williams (1930); Larsen (1975, 1984, 1986) | Larsen (1987) |
| <i>Cupido argiades</i> | Williams (1938); Williams <i>et al.</i> (1942); Baker (1969); Vermandel & Vliegenthart (2014) | Parmesan <i>et al.</i> (1999) |
| <i>Danaus chrysippus</i> | Evershed (1910); Williams (1930); Williams (1938); Dell (1977); Larsen (1984); Braby (1988); Harris (1988); Dingle <i>et al.</i> (1999); Smith (2014) | Bhaumik & Kunte (2018) |
| <i>Danaus gilippus</i> | Williams (1930); Miller (1985); Simon & Miller (1986); Gibo (1993); Haber (1993) | Brower (1995) |
| <i>Euphydryas editha</i> | Harrison (1989) | Gilbert & Singer (1975) |
| <i>Euploea midamus</i> | Williams (1930); Chan (2004); Wong <i>et al.</i> (2004); Yiu (2009) | Bhaumik & Kunte (2018) |
| <i>Euploea mulciber</i> | Williams (1930); Wang & Emmel (1990); Orr (1992); Chan (2004); Wong <i>et al.</i> (2004); Yiu (2009) | Bhaumik & Kunte (2018) |
| <i>Eurema herla</i> | Williams (1930) | Braby (2005a) |
| <i>Eurema laeta</i> | Williams (1927, 1930, 1938) | Braby (2005a) |
| <i>Heliconius ethilla</i> | Williams (1930); Beebe (1950a) | Ehrlich & Gilbert (1973) |
| <i>Maniola jurtina</i> | Williams <i>et al.</i> (1942) Baker (1969) | Parmesan <i>et al.</i> (1999) |
| <i>Nymphalis californica</i> | Williams (1930); Shapiro (1982, 1986) | Shapiro (1976) |
| <i>Parnassius apollo</i> | Nakonieczny <i>et al.</i> (2007) | Parmesan <i>et al.</i> (1999) |
| <i>Parnassius mnemosyne</i> | Välímäki & Itämies (2003); Vlasanek <i>et al.</i> (2009); Bolotov <i>et al.</i> (2013) | Parmesan <i>et al.</i> (1999) |

| Species | Recorded as migratory | Recorded as non-migratory |
|-------------------------------|--|-------------------------------|
| <i>Polygonia c-album</i> | Baker (1969); Beshkov (1996a, b) | Parmesan <i>et al.</i> (1999) |
| <i>Pyronia tithonus</i> | Cook <i>et al.</i> (2001) | Parmesan <i>et al.</i> (1999) |
| <i>Tirumala septentrionis</i> | Williams (1927); Evershed (1910); Fisher (1945); Chan (2004); Larsen (1987, 2004); Yiu (2009); Ramesh <i>et al.</i> (2012) | Bhaumik & Kunte (2018) |

VI. STUDYING BUTTERFLY MIGRATION

Migration syndromes were first recorded in the early–19th century by Charles Darwin during his famous journey on the Beagle (Darwin, 1903), followed in the early 20th century by other notable contributions (Hayward, 1925, 1953, 1967; Williams, 1957, 1958). Since then, observing and recording migrating butterflies has become popular among professional and amateur lepidopterists worldwide, and now hundreds of species are known to be migratory (Fig. 4).

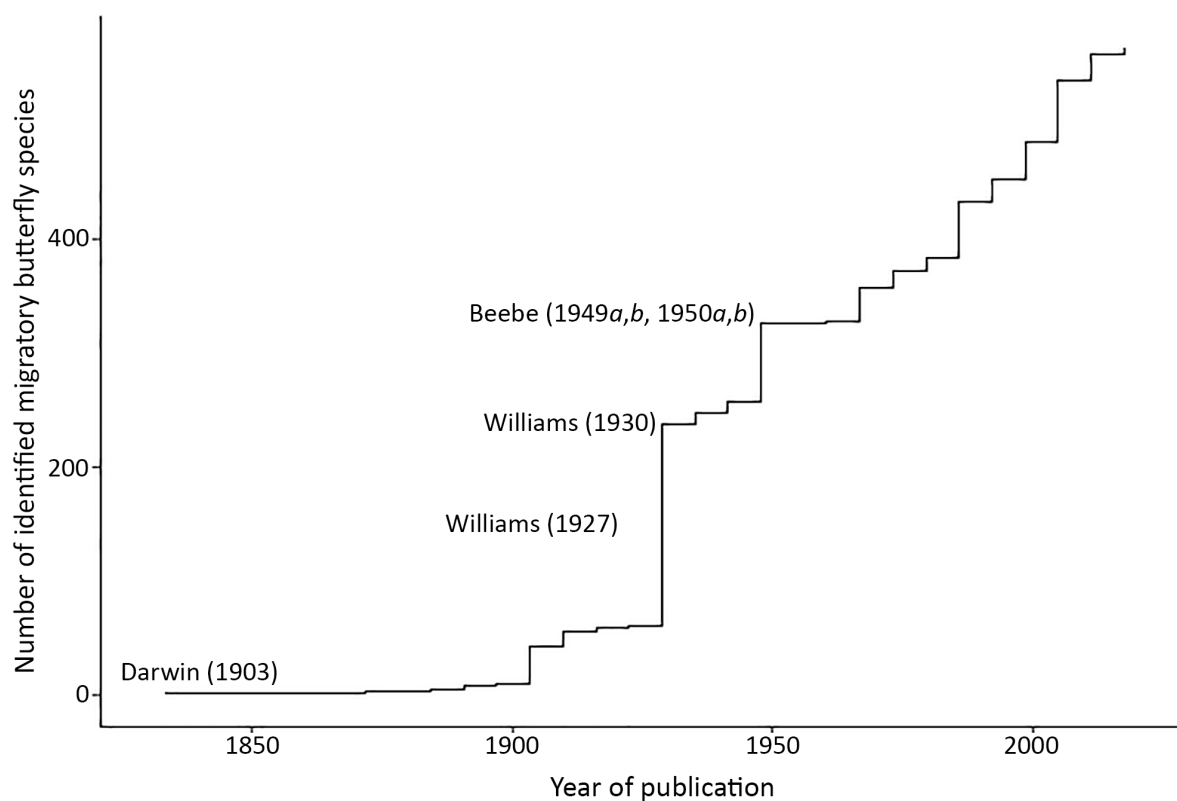


Fig. 4. Cumulative number of butterfly species documented as being migratory. Key publications responsible for rapid increases in the list of known migrants are noted.

Discoveries of migratory butterflies have increased steadily in recent decades (Fig. 4), yet much of this research effort is focused on a very small number of species (notably *Danaus plexippus*, *Vanessa cardui*, and *Vanessa atalanta*) with little work on the majority of species, substantially biasing knowledge of migration among butterflies globally. Nearly 55% of the studies we located were focused on just two species, *D. plexippus* (215 studies) and *V. cardui* (124 studies). We found only one or two research studies for 76% of migratory butterflies, with more than 10 publications for only 5% of butterflies recorded as migratory. For less-studied species, most authors used relatively weak lines of evidence to diagnose migration, such as records of mass movement, or nomadic behaviour. Very few studies have reported physiological (5%), and experimental (11%) lines of evidence (Fig. 1).

Thaxter (1880) was the first to describe the overwintering sites of various butterflies including monarchs, along the California coast. However, it took nearly a century to elucidate the monarch migration cycle fully (Malcolm *et al.*, 1993; Brower, 1995) and to identify their major overwintering site in the Transvolcanic Mountains of central Mexico following nearly 38 years of study, including 24 years of tagging studies (Urquhart & Urquhart, 1976). Until the 1960s, most research on butterfly migration was experimental, but in the early 1960s, C.G. Johnson investigated differences in life-history traits between migratory and non-migratory butterflies. Johnson found that migratory females start their flight before ovarian development (i.e. they are sexually immature), that they usually increase in body mass prior to migration, and that they possess longer forewings (Johnson, 1963, 1969). Genetic techniques were first used to study migration in the mid-1970s, when Eanes & Koehn (1978)

collected summer migrant monarchs from the USA and Canada, and used electrophoretic alleles at six enzyme loci as genetic markers to study the genetic structure of the population. They found significant allele frequency variation between migratory and non-migratory individuals (Eanes & Koehn, 1978). Malcolm *et al.* (1993) used the presence of plant-derived cardenolides in monarch butterflies to describe spring migration in relation to the temporal and spatial distribution of about 130 species of their larval host plant, milkweed (*Asclepias* spp.). In recent decades, there has been rapid improvement in the use of genetic tools to uncover patterns of migratory connectivity (Sauman *et al.*, 2005; Dingle, 2006; Zhu, Casselman & Reppert, 2008; Zhan *et al.*, 2014; Reppert & de Roode, 2018), specialised radar to detect high-flying migratory butterflies (Chapman *et al.*, 2010; Stefanescu *et al.*, 2013; Chapman & Drake, 2016), isotopic tools to track the origin of migratory butterflies (Wassenaar & Hobson, 1998; Flockhart *et al.*, 2013, 2015), and tethered flight studies to identify migratory orientation (Mouritsen & Frost, 2002; Reppert, Zhu & White, 2004; Nesbit *et al.*, 2009).

VII. CONSERVING MIGRATORY BUTTERFLIES

Migratory species can be acutely vulnerable to threats, since they depend on a series of habitats throughout their migration route (Runge *et al.*, 2014). Yet, reports on the conservation status of migratory butterflies are sparse except for monarchs (Brower *et al.*, 2012; Flockhart *et al.*, 2015; Semmens *et al.*, 2016; Agrawal & Inamine, 2018; Agrawal, 2019). While European populations of some well-known migratory butterflies are considered stable [e.g. painted lady *V. cardui*, red admiral *V. atalanta*, and clouded yellow *Colias croceus* (Fox *et al.*, 2015)], in North America there is very good evidence of a strongly declining population trend in the monarch.

In North America, the monarch butterfly is declining so rapidly (80% over the last decade) that it may soon lead to the demise of this world-famous migration (Brower & Malcolm, 1991; Semmens *et al.*, 2016). Several factors including loss of larval food plants (milkweed *Asclepias* spp.) and nectar resources, degraded overwintering habitats, climate change, influence of invasive milkweeds and natural enemies, global wind patterns, artificial light, habitat fragmentation, pathogens, agriculture and genetically modified crops, and declining fecundity, etc., have all been implicated, demonstrating the complexity of diagnosing and reversing declines in populations of migratory species (Malcolm *et al.*, 1993; Brower *et al.*, 2012; Oberhauser, Nail & Altizer, 2015; Oberhauser *et al.*, 2017; Pleasants *et al.*, 2016, 2017; Stenoien *et al.*, 2016; Thogmartin *et al.*, 2017; Malcolm, 2018; Agrawal, 2019; Freedman *et al.*, 2020).

The abundance and diversity of butterflies is controlled to a great extent by the presence and absence of food plants (Zalucki & Suzuki, 1987; Brower *et al.*, 2012; Batalden, Oberhauser & Peterson, 2014; Robertson, Zalucki & Paine, 2015; Pleasants & Oberhauser, 2017; Chowdhury *et al.*, 2020, 2021), and a landscape-level reduction of host plants can cause huge impacts on migrants (Zalucki & Kitching, 1982; Zalucki & Suzuki, 1987; de Roode *et al.*, 2008; Dyer & Forister, 2016; Pleasants & Oberhauser, 2017). For example, in North America, the use of genetically modified crops can impact insect-herbivore relationships and interspecific competition with other weeds (Rehfeldt *et al.*, 2012). The availability of herbicide-tolerant crops has changed agricultural practices from conventional tillage to no-till, and milkweed has been extirpated in soy and corn fields by the use of large quantities of glyphosate (Zalucki, Brower & Alonso, 2001). Genetically modified maize plants expressing crystalloproteins from the soil bacterium *Bacillus thuringiensis* (Bt) that are toxic to lepidopteran larvae can increase monarch larval mortality by 44% (Losey, Rayor & Carter,

1999). Field experiments showed that Bt maize pollen deposited on potted plants is toxic to first-instar monarch larvae (Jesse & Obrycki, 2000). Currently, the general consensus is that the risk to monarchs from Bt pollen is low because the fraction of the population exposed is small (Sears *et al.*, 2001).

Concern is increasing regarding the use of neurotoxic neonicotinoids, such as imidacloprid seed treatments, to control arthropod pests (Malcolm, 2018). The insecticide is present in the pollen and nectar of treated plants (Cresswell, 2011), leading to lethal to sub-lethal doses of imidacloprid being taken up by nectar-feeding butterflies (Stenoien *et al.*, 2016; Malcolm, 2018). Smoke from agricultural fires near the Mexican overwintering site for monarchs, which drifts up the mountainsides into the colonies of hibernating butterflies, is considered an ominous threat (Brower & Malcolm, 1991). When the air temperature is near the flight threshold, smoke causes frenzied flight, but if the temperature is below this threshold, the butterflies drop to the ground with a substantially increased predation risk (Brower, 1977; Calvert, Hedrick & Brower, 1979; Glendinning, Mejia & Brower, 1988; Brower & Malcolm, 1991).

The availability of nectar plants also depends on precipitation; thus, drought will influence the presence of nectar for migratory butterflies such as monarchs (Stenoien *et al.*, 2016; Thogmartin *et al.*, 2017). For example, monarchs feeding on irrigated *Liatris mucronata* were able to store up to 80mg of lipids in their body tissues, whereas those feeding on the drought-stressed *Verbesina virginica* plants could only store half this amount (Brower *et al.*, 2015). In addition, fire suppression can play a key role in the abundance of nectar resources (Rudolph *et al.*, 2006).

Habitat loss to urbanisation and agricultural expansion reduces stopover site availability, which will eventually lead to higher densities of migrants using fewer sites (Wilcove &

Wikelski, 2008; Flockhart *et al.*, 2013, 2015; Chowdhury *et al.*, 2017). In general, high local densities and greater species diversity can lead to increases in both between- and within-species transmission of pathogens (Bradley & Altizer, 2005; Altizer, Bartel & Han, 2011; Satterfield, Maerz & Altizer, 2015).

Butterflies are sensitive to environmental factors, and slight changes can cause rapid population declines. Climate change is likely to affect moisture and temperature regimes, and thereby disrupt migratory cues. Anderson & Brower (1993) experimentally determined that surface water on the bodies of monarch butterflies at overwintering sites in Mexico affects their survival at low temperatures: 50% of dry butterflies were killed at -8°C , and 100% at -14°C , but if their wings and bodies were wet, 50% died at -4°C , and 100% at -8°C . Larsen & Lee (1994) showed that wet monarchs freeze more rapidly than dry individuals and discussed the importance of intact oyamel fir forest to their survival. In addition, migrants are affected by wind patterns: stronger winds can carry the migrants far from their destination, and weak winds may increase the energy required to reach the destination (Cardé, 2008; Chapman *et al.*, 2008; Sydeman *et al.*, 2014).

Temperature and the availability of water resources can influence the abundance of migratory butterflies in various ways (Brower, Fink & Walford, 2006; Robinson *et al.*, 2009; Chowdhury *et al.*, 2021). Temperature and rainfall patterns are changing rapidly, and up to 69% of the overwintering area of the monarch butterfly could become unsuitable by 2030, with possibly none remaining suitable by the end of this century (Sáenz-Romero *et al.*, 2012; Ramírez *et al.*, 2015). Climate change is also associated with increased winter storms in Mexico, leading to substantial mortality of overwintering monarchs (Oberhauser & Peterson, 2003; Brower *et al.*, 2004; Berry *et al.*, 2005; Batalden *et al.*, 2014). Milkweeds are projected to shift northward in North America in both moderate ($1-3^{\circ}\text{C}$ increase) and severe ($2-6^{\circ}\text{C}$

increase) climate warming scenarios, with the potential for environmental cues for migration (e.g. photoperiod) becoming decoupled from the availability of food plants (Lemoine, 2015; Borkin, 2017).

The directional movement of migratory butterflies means they must cross linear features such as roads, and collisions with vehicles can cause substantial mortality (Kantola *et al.*, 2019; Tracy *et al.*, 2019). It has been estimated that 20 million butterflies including 500,000 monarchs are killed along roads in Illinois each week (McKenna *et al.*, 2001; Berenbaum, 2015). Kantola *et al.* (2019) estimated that nearly 2.1 million monarchs are killed annually in the Central Funnel in Texas, representing around 3% of the overwintering population. The presence of milkweed larval host plant (*Asclepias verticillata*) and other nectar resources (e.g. Asteraceae, *Trifolium* spp., and Apiaceae) along road verges means that roads attract butterflies, and measures such as speed-limiting vehicles at roadkill hotspots, especially during the peak migration period, could substantially reduce mortality (Kantola *et al.*, 2019). Obtaining credible data on the survival rates of butterflies during migration is difficult (Flockhart *et al.*, 2015), but such information will be essential for assessing the success of conservation interventions.

The causes of declines in migratory butterflies can be very difficult to diagnose. In the case of the monarch, much attention has focused on threats to their overwintering sites in Mexico (Brower *et al.*, 2002, 2012; Oberhauser & Peterson, 2003; Ramírez, Azcárate & Luna, 2003; Agrawal & Inamine, 2018), but as detailed above there are also substantial threats in their breeding range and during migration. It is likely that climate change (especially extreme events), and the distribution of host plants represent the most significant threats to migratory butterflies (Marini & Zalucki, 2017; Malcolm, 2018; Chowdhury *et al.*, 2020, 2021).

We have focused our discussion in this section on just one particularly well-studied migratory butterfly species, which is rather atypical in some ways: for example, its mixed strategy of migration to adult hibernation refugia is very unusual (perhaps unique) among migrant butterflies. By contrast, for the many species of ‘more typical’ butterfly migrants, there is a real lack of knowledge about population trends and conservation status. One of the few relatively well-studied species is the painted lady; this species carries out long-range migrations of approximately the same distance as monarchs (Stefanescu *et al.*, 2013), and also feeds on fast-growing weedy host plants of ephemeral habitats. However, in contrast to monarchs, painted ladies do not appear to be declining, at least in Europe (Fox *et al.*, 2015). A comparative study investigating similarities and differences between these two migrant species, and the impact of various life-history traits on their population trends would be instructive. Further work on the factors influencing decline in migratory butterflies, especially population-level studies, is urgently needed to assess connectivity between breeding and non-breeding grounds (Gao *et al.*, 2020) and population dynamics across the whole annual cycle.

VIII. CONCLUSIONS

(1) Nearly 600 butterfly species are currently known to be migratory, and the rate at which new migrants are discovered shows no sign of slowing down (Fig. 4). Migratory butterflies occur all over the world except Antarctica, with most records from tropical countries with wet-dry seasonality (e.g. Australia, India, and Sri Lanka). However, the largest proportions of butterflies identified as migrants are from relatively well-studied regions of the world such as Europe, USA, and Australia, probably reflecting their greater interest in research on butterflies.

(2) We found 24 cases where a species was noted as sedentary by some authors, but was considered a migrant elsewhere in its range. This might reflect errors of course, but perhaps more likely suggests that such species have both migrant and non-migrant subgroups within the same population (partial migration; Menz *et al.*, 2019). Using stable isotope analyses, Van der Zanden *et al.* (2018) recently found that 48% of monarchs from South Florida are migrants while the remainder are residents, lending support to the idea that partial migration might be common in butterflies.

(3) The proportion of migratory butterflies varied enormously taxonomically. Some 53% of Libytheinae butterflies are migratory, while the proportion is much lower in other subfamilies. Future studies should obtain regional records, document movement patterns, and conduct experimental work on migration in butterflies to reveal hotspots and their vulnerability to extinction, thus providing invaluable data for butterfly conservation. A fruitful avenue would be to focus on mapping the distributions of migratory butterflies at different times of year.

(4) Many butterflies migrate from one part of the world to another every year. They cross international borders, and the biomass transfer during migratory flight is extensive (Hu *et al.*, 2016). Despite this, we possess limited knowledge on their origin, stopover sites, and destination (Pierce *et al.*, 2014). It took decades for scientists to establish the importance of studying butterfly migration and newer methods to investigate migratory patterns are now available. Although multiple methods to describe migratory behaviour exist, applications of most of these has been limited to just a few species (e.g. *D. plexippus*, *V. cardui*, *V. atalanta*). We remain far from understanding the true prevalence of migration in butterflies, and the existing evidence is rather weak for many species.

(5) About half of all lepidopteran species are estimated to be decreasing in abundance worldwide (Sánchez–Bayo & Wyckhuys, 2019; van Strien *et al.*, 2019; Van Klink *et al.*, 2020). If the acute declines recorded for the North American monarch are replicated in other migratory butterflies around the world, there might be hundreds of such declines that are going undetected. Successful conservation of migratory butterflies will depend on achieving a much better understanding of which species are migratory, and whether and why populations are declining.

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X. SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of the article.

Table S1. Species determined to be migratory based on studies using a variety of different experimental, physiological or population evidence methods.

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