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Dispersal strategies in terrestrial insects Mark K Asplen

- 4 Terrestrial insects frequently disperse and/or migrate, either
- 5 through their own self-directed actions or via other vehicles.
- 6 Here, the following recent advances in the study of insect
- 7 dispersal are highlighted: (1) components of classic
- 8 hypotheses (marginal value theorem and inbreeding avoidance
- 9 via sex-specific dispersal) have found varying degrees of
- ¹⁰ recent support; (2) modern genetic tools have uncovered
- several candidate dispersal genes; (3) dispersal syndromes
- vary in their genetic and/or physiological constraints; and (4)
- 13 common laboratory techniques may not accurately reflect
- dispersal in the field. A common theme is the tendency for
- ¹⁵ breakthroughs to be concentrated in species with extremely
- 16 well-defined dispersal phenotypes (e.g., long-distance
- 17 migrants, wing polymorphic insects), suggesting the need for
- 18 increased focus on species exhibiting less self-directed modes
- 19 of dispersal.

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27 Introduction

Class Insecta is known for two of the greatest evolutionary 28 milestones in Animalia: (1) it contains, by far, the largest 29 number of species (approximately 5.5 million) [1]; and (2) 30 it is the first lineage to have evolved powered flight 31 (possibly during the Early Devonian) [2]. An evolutionary 32 advantage of the latter is increased capacity for dispersal 33 and/or migration (see Box 1 for discussion of the distinc-34 tion between the two terms), which can yield substantial 35 adaptive benefits such as avoidance of environmental 36 stressors and access to seasonally-available habitats [3]. 37 Dispersal is, of course, also critical to the study of extant 38 insect populations via its importance in predicting their 39 establishment, seasonality of occurrence, and rate of 40 spread [4-6]. 41

As with all behaviors, there is value to studying both the 42 proximate and ultimate causes of insect dispersal strate-43 gies (Figure 1). This brief review utilizes this approach to 44 highlight recent advances, with specific reference to (1) 45 classic hypotheses of how dispersal may be optimized via 46 natural selection; (2) the genetics behind dispersal, which 47 determines its evolutionary potential; and (3) the evolu-48 tion of suites of life history traits ('syndromes') correlated 49 with dispersal by flight. Finally, I will address the more 50 practical issue of whether or not laboratory techniques 51 commonly employed by entomologists accurately reflect 52 insect dispersal in the field. 53

Optimization of dispersal decisions in insects 54

Optimality models of dispersal commonly emphasize the 55 following benefits of leaving a site, weighed against the 56 safety risks [7]: (1) increased habitat sampling, and (2) 57 avoidance of inbreeding depression at the natal site. With 58 respect to the former, the marginal value theorem (MVT) 59 posits that an optimal forager adjusts patch residency time 60 based on resource richness and time costs [8,9]. For the 61 MVT to be realistic, however, insects must be able to 62 gather information on these variables. Although host 63 location cues have received considerable attention in 64 insects, recent studies of parasitoid wasps by Parent 65 and colleagues [10[•],11[•]] have confirmed their ability to 66 measure time while foraging. First, the egg parasitoid 67 Trichogramma euproctidis appears to measure time by remembering the duration of the pre-oviposition host 68 assessment period (the 'initial transit duration') [10[•]]. 69 Female T. euproctidis lay more eggs on hosts for which 70 the initial transit duration was longer, an effect that is 71 independent of host egg size. Another parasitoid species, 72 the braconid *Microplitis croceipes*, is able to associate time 73 intervals with odors [11[•]]. In this study, female parasitoids 74

Box 1 The distinction between dispersal and migration. Dispersal and migration are related concepts that often have similar effects on gene flow in populations. Although recognizing the myriad definitions and population-level outcomes of migration [40], migration is defined here as a characteristic of individuals and contains the key attributes noted by Kennedy [41]: (1) inhibition of responses to typical 'station keeping' cues used to locate resources; (2) abandonment of the home range via this extended movement; and (3) as a behavior, it is 'persistent and straightened out', occurring regardless of whether the animal is using its own locomotory apparatus or not (e.g., using a current). Dispersal has been defined in two ways [40], the latter of which is emphasized in this review: (1) an increase in distance between members of a population: or (2) behaviors leading to displacement from the natal habitat or movements between breeding habitats. Here, it is recognized that migration produces movements that may (or may not) result in dispersal according to each sense of the word.

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2 Behavioral ecology

Figure 1



Dispersal and migratory behaviors have both proximate and ultimate causes. Proximate mechanisms include a behavior's genetic basis, as well as its potential regulation through physiological (neuronal, hormonal, or cell signaling pathways) or developmental processes (e. g., learning). Ultimate mechanisms explain the evolution of a behavior, both in terms of optimization via adaptation and evolutionary constraints due to life history trade-offs or phylogenetic history. The two migratory insect species pictured differ greatly in the degree to which we understand this aspect of their biology, illustrating the need for more research on non-traditional insect models (see text). Top: North American populations of the monarch butterfly (Danaus plexippus) exhibit seasonal, transgenerational migrations on a continental scale [42]. There are clear signs of a dispersal syndrome in this species: migratory individuals possess significantly larger forewings [27], and reproductive diapause is coupled with increased longevity (modulated by juvenile hormone) during the migratory period [43]. Furthermore, the molecular and genetic basis of monarch migration has been the subject of extensive study [e.g., 41]. Bottom: The sweet potato whitefly (Bemisia tabaci) exhibits migratory behaviors during vertical flight chamber assays, with approximately 6% of the population ignoring a green light vegetative cue in favor of sustained, positively phototactic climbs [45]. The genetic basis of this behavior has not been studied, and the species appears to exhibit only weak signs of a dispersal syndrome (i.e., subtle differences in wing morphology and mixed evidence for an oogenesis-flight syndrome) [45-47].

were trained by offering them caterpillar hosts after 5 min 75 (short) or 30 min (long) intervals, each associated with a 76 different odor cue (strawberry or vanilla). Successful 77 conditioning was measured using a 'choice test' of trained 78 parasitoids placed downwind of both odor sources follow-79 ing either a short or long interval since host exposure. 80 Wasps in this assay consistently preferred the correct 81 pairing of time interval experienced and its associated 82 odor, with no biases connected to chamber position of the 83 odor, training-test order, or the odor itself. Interestingly, 84 restraining the parasitoid during training interferes with 85 this process, suggesting that energy expenditure may also 86 play a role in time measurement. 87

A common trend in insects is the tendency for one sex to exhibit higher dispersal rates [12,13]. Such sex-biased dispersal has long been considered an adaptation for inbreeding avoidance [14], especially in hymenopterans with complementary sex determination (such as all known members of Formicidae) [15–17]. Here, low fitness diploid males (rather than normal haploid males) 94 can be produced from fertilized eggs if homozygosity 95 (which is expected to increase with inbreeding) exists 96 at one or more sex determination loci. An extensive recent 97 study with the ant *Formica exsecta* suggests that, despite 98 substantial evidence of male-biased dispersal (twice the 99 distance on average of queens), neither male nor female 100 natal dispersal distance correlated with a reduction of 101 homozygosity in workers [18]. Detailed studies that thor-102 oughly test long-held hypotheses under field conditions 103 are important for understanding the adaptive basis of sex-104 biased dispersal in insects, which may also have maladap-105 tive consequences (e.g., increased risks of Allee effects) 106 [5,19]. 107

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The genetic basis of insect dispersal strategies

Although the most famous candidate 'dispersal gene' is 110 foraging (for), with its 'rover-sitter' polymorphism first discovered in Drosophila melanogaster [20], there have 111 been several recent studies searching for additional ones. 112 Perhaps the most pivotal of these combined measure-113 ments of tethered flight (i.e., behaviors measured while 114 the insect is attached to a flight mill; see the section on 115 laboratory measures of dispersal below) with genomic 116 analysis, revealing several candidate genes for migratory 117 activity in the cotton bollworm (*Helicoverpa armigera*) 118 [21^{••}]. In this species, short and long-distance flight 119 activity is associated with the differential expression of 120 approximately 1.26% of the genome, with a considerable 121 proportion of the 215 genes associated with physiological 122 factors classically related to insect migration: lipid mobi-123 lization (presumably for flight fuel, although not all insect 124 taxa use lipids for this purpose [22]), reduction of oxida-125 tive stress in highly active tissues, flight muscle structure, 126 and juvenile hormone regulation. Another approach 127 involves direct genetic manipulation to study flight-128 related traits. In one example, RNA interference in the 129 nub-Gal4 gene of D. melanogaster can produce wing shapes that increase in-flight maneuverability in laboratory are-130 nas [23]. Perhaps more germane to long-distance move-131 ment, the gene editing technologies TALENs and 132 CRISPR/Cas9 have recently been employed in the mon-133 arch butterfly (Danaus plexippus). Here, mutagenesis of 134 the *clock* gene disrupts the circadian clockwork of this 135 species, which is predicted to have strong effects on this 136 seasonal migrant [24]. 137

The recent advances outlined above are critical steps 138 toward understanding the genetic architecture of insect 139 dispersal, especially in long-distance migrants with well-140 defined phenotypes (see the discussion of dispersal syn-141 dromes). Terrestrial insect species vary greatly, however, 142 in the degree to which they control potentially substantial 143 movements through the environment. This certainly 144 limits our ability to generalize from these important 145 case studies. Some species disperse passively through 146

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