

Model Systems in Ecology, Evolution, and Behavior: A Call for Diversity in Our Model Systems and Discipline*

Meghan A. Duffy,^{1,†} Carlos García-Robledo,² Swanne P. Gordon,³ Nkrumah A. Grant,⁴ Delbert A. Green II,¹ Ambika Kamath,⁵ Rachel M. Penczykowski,³ María Rebollo-Gómez,⁶ Nina Wale,⁷ and Luis Zaman^{1,8}

1. Department of Ecology and Evolutionary Biology, University of Michigan, Ann Arbor, Michigan 48109; 2. Department of Ecology and Evolutionary Biology, University of Connecticut, Storrs, Connecticut 06269; 3. Department of Biology, Washington University in St. Louis, Saint Louis, Missouri 63130; 4. Department of Biological Sciences, University of Idaho, Moscow, Idaho 83844; 5. Miller Institute for Basic Research in Science, University of California, Berkeley, California 94720; 6. Department of Ecology and Evolutionary Biology, Yale University, New Haven, Connecticut 06520; 7. Department of Microbiology and Molecular Genetics, Department of Integrative Biology, and Ecology, Evolution, and Behavior Program, Michigan State University, East Lansing, Michigan 48824; 8. Center for the Study of Complex Systems, University of Michigan, Ann Arbor, Michigan 48109

Submitted February 15, 2021; Accepted March 26, 2021; Electronically published May 21, 2021

Online enhancements: supplemental table.

ABSTRACT: Ecologists and evolutionary biologists are fascinated by life's variation but also seek to understand phenomena and mechanisms that apply broadly across taxa. Model systems can help us extract generalities from amid all the wondrous diversity, but only if we choose and develop them carefully, use them wisely, and have a range of model systems from which to choose. In this introduction to the Special Feature on Model Systems in Ecology, Evolution, and Behavior (EEB), we begin by grappling with the question, What *is* a model system? We then explore where our model systems come from, in terms of the skills and other attributes required to develop them and the historical biases that influence traditional model systems in EEB. We emphasize the importance of communities of scientists in the success of model systems—narrow scientific communities can restrict the model organisms themselves. We also consider how our discipline was built around one type of “model scientist”—a history still reflected in the field. This lack of diversity in EEB is unjust and also narrows the field's perspective, including by restricting the questions asked and talents used to answer them. Increasing diversity, equity, and inclusion

will require acting at many levels, including structural changes. Diversity in EEB, in both model systems and the scientists who use them, strengthens our discipline.

Keywords: model organisms, biodiversity, diversity in STEM, equity, inclusion, model system.

“What is true for *E. coli* is true for the Elephant”
—J. Monod

“But not for *Salmonella*” —E. Groisman
(Aisha Burton, Twitter post, January 14, 2021,
9:48 a.m.)

Introduction

As scientists studying ecology, evolutionary biology, and behavior, we love, celebrate, and are captivated by life's diversity—those “endless forms most beautiful,” as Charles Darwin famously framed it. At the same time, we seek to understand how the natural world works—to identify general phenomena and the mechanisms driving them. Indeed, the American Society of Naturalists has identified “conceptual unification of the biological sciences” as its purpose. Thus, ecologists and evolutionary biologists face a challenge: extracting general principles and mechanisms from amid all the wonderful diversity surrounding us (Kokko 2020). We aim to see the forest *and* the trees.

Model systems can help us meet this conceptual unification-despite-abundant-diversity challenge (Kokko 2020), but only if we choose (and develop) our model systems carefully, use them wisely, and have a range of model

* This article originated as part of the 2020 Vice Presidential Symposium. The symposium was originally scheduled for the 2020 annual meeting of the American Society of Naturalists in Cleveland, Ohio; because of the COVID-19 pandemic, it instead occurred at the Virtual Asilomar meeting in January 2021.

† Corresponding author; email: duffymeg@umich.edu.

ORCIDs: Duffy, <https://orcid.org/0000-0002-8142-0802>; García-Robledo, <https://orcid.org/0000-0002-5112-4332>; Gordon, <https://orcid.org/0000-0002-9840-725X>; Grant, <https://orcid.org/0000-0002-4555-5283>; Green, <https://orcid.org/0000-0002-3373-8867>; Kamath, <https://orcid.org/0000-0002-4012-0483>; Penczykowski, <https://orcid.org/0000-0003-4559-0609>; Rebollo-Gómez, <https://orcid.org/0000-0002-3592-4479>; Wale, <https://orcid.org/0000-0001-9703-444X>; Zaman, <https://orcid.org/0000-0001-6838-7385>.

systems from which to choose. Despite what Jacques Monod claimed, what is true for *Escherichia coli* is not necessarily true for the elephant. Indeed, even if we consider things at a narrower scale, mice, zebra fish, *Caenorhabditis elegans*, and fruit flies cannot represent all animals; *Arabidopsis* cannot represent all plants; and *E. coli* and *Saccharomyces cerevisiae* cannot represent all microbes.

The choice of study organism (or system)—and its match to the question under study—is critical to our science (Travis 2006). Researchers consider myriad factors when choosing a study organism (Dietrich et al. 2020). Choice of study organism is often influenced by tractability (Krogh 1929; Green et al. 2018) and also reflects the impact of access, resources, and economies (Burian 1993; Dietrich et al. 2020)—which means that there are biases in our current model systems, as we discuss more below. Study organisms are also chosen because they might enable comparisons to other organisms, which can reveal general phenomena and processes (Burian 1993; Travis 2006; Dietrich et al. 2020). The knowledge we collectively build reflects thousands of individual decisions regarding which systems should be used to study which questions (Travis 2006); however, these decisions are not fully independent but, rather, are influenced by social networks, prior research, mentoring relationships, and other factors. In the end, “the principles and facts that emerge will be only as reliable as our choices have been wise” (Travis 2006, p. 303).

Ecologists and evolutionary biologists need a diversity of study systems to achieve our goal of conceptual unification, and we must be thoughtful and creative about how we use and develop those systems. This special feature highlights a variety of ways in which model systems are currently being used to address timely and important questions in ecology and evolutionary biology (García-Robledo and Baer 2021; Gordon et al. 2021; Grant et al. 2021; Green 2021; Penczykowski and Sieg 2021; Wale and Duffy 2021). In this introduction, we first seek to define what we mean by the term “model system” (a surprisingly challenging task). Having done that, we then ask where our model systems come from (both in terms of the skills required to develop them or use them in new ways and in terms of their history and geography), and we also consider where model systems research in ecology, evolution, and behavior (EEB) might be heading.

In an article addressing the importance of diversity in ecology and evolutionary biology, it is essential to emphasize that EEB needs diversity not just in terms of what organisms we study but also in terms of who does those studies. EEB as a field was built around one type of “model scientist”—someone who is white, male, cisgendered, affluent, not disabled, and without major caregiving responsibilities (to list only a few salient features). Unfortunately, the demographics of our field still reflect those origins (Rushworth et al. 2021), as do science, technology, engineering, and mathematics (STEM) more broadly (McGee 2020). This lack of

representation in EEB is a clear moral and ethical issue, which on its own makes this important to address. In addition, as we discuss more below, this lack of diversity narrows the field’s perspective and holds back our science. Thus, we also discuss some of the impacts of the lack of diversity in EEB and cover strategies for achieving a more diverse, equitable, and inclusive discipline. Diverse scientists will yield the diverse model systems and diverse perspectives that EEB needs if we are to meet the challenge of identifying the general principles and mechanisms that generate endless forms most beautiful.

What Is a Model System?

“Model organism,” “model species,” and related terms have been criticized as some of the most overused and underdefined words in life sciences (Katz 2016). It is therefore with some trepidation that we seek to define what a model system is for our purposes.

In biomolecular sciences, model organisms are experimental organisms that are studied in the laboratory context as representatives of a broad range of organisms and processes (Jenner and Wills 2007; Ankeny and Leonelli 2011; Leonelli and Ankeny 2013); in biomedical sciences, model organisms are often chosen (and developed) on the basis of similarity to humans. Classical model organisms often exhibit a number of characteristics that make them amenable to laboratory life, including short generation times, small size, and ease of manipulation and measurement, which is why Bolker (1995) argued that “model systems are likely to be peculiar with respect to their own taxa, but relatively consistent with respect to each other” (p. 451). Biomolecular researchers who work on model organisms tend to share the rationale that despite the (unusual) biological characteristics that make model organisms models, the conclusions one draws from them are generalizable because traits are evolutionarily conserved (Ankeny and Leonelli 2011, 2020), and that understanding core biomolecular phenomena is best achieved by divorcing organisms from their ecological context (Ankeny and Leonelli 2020). The processes used to construct these “traditional” model organisms (including standardization and modes of manipulation) and the scientific culture that surrounds their study are as much a part of what makes an organism a model as are their inherent biological traits (Ankeny and Leonelli 2020).

There are challenges to adopting the biomolecular definition of a model system for EEB because our fundamental goal is different. We seek to understand genetic and phenotypic *variation* and how the context in which organisms live modulates this variation. As such, the organism’s environment is a feature of the system that must also be studied (Bartholomew 1966)—some of us would even argue that there is no meaningful organism without its environment (e.g., Lewontin 2001).

Here, we propose the following definition of what a model system is for EEB: a species, taxon, community, or ecosystem that has been studied from multiple angles with a goal of developing a deep understanding of that organism (or taxon, community, or ecosystem), in a manner that enables comparisons with other systems to illuminate general ecological, evolutionary, and/or behavioral principles; achieving this will require that the system has been studied long enough for a substantive body of knowledge to have been generated. Model systems are designated as such by the community—a single person cannot decide on their own that something is a model system; crucially, this can lead to gatekeeping and adds to the importance of having diversity in our discipline, as we discuss more in the second half of this article. Prior discussions of model systems in EEB have contrasted work on model systems versus natural populations (Travis 2006). However, the terms “model system” and “natural population” are not mutually exclusive. Rather, we propose that there are multiple axes along which

model systems fall (fig. 1). We are in full agreement with Travis (2006) that “robust inference requires horizontal comparisons and vertical integration” (p. 307)—the first part of our definition is Travis’s vertical integration (i.e., the study within a single system of processes at a number of levels of biological organization), and the second part is his horizontal comparisons (i.e., when a single question is studied at the same level of biological organization in multiple systems). In our definition, a model system need not necessarily be a single species (or taxon) or a pair of closely interacting species. Moreover, it need not be amenable to laboratory study. Instead, in EEB certain sites and ecosystems have also emerged as model systems as a result of an extended history of study that has allowed us to generate and test general ecological and evolutionary theory (table S1, available online; fig. 1). This includes experimental species assemblages, such as the Cedar Creek biodiversity plots; natural tree plots, such as the Forest Census Plot on Barro Colorado Island; and networks of such ecosystems, such as the 72-site

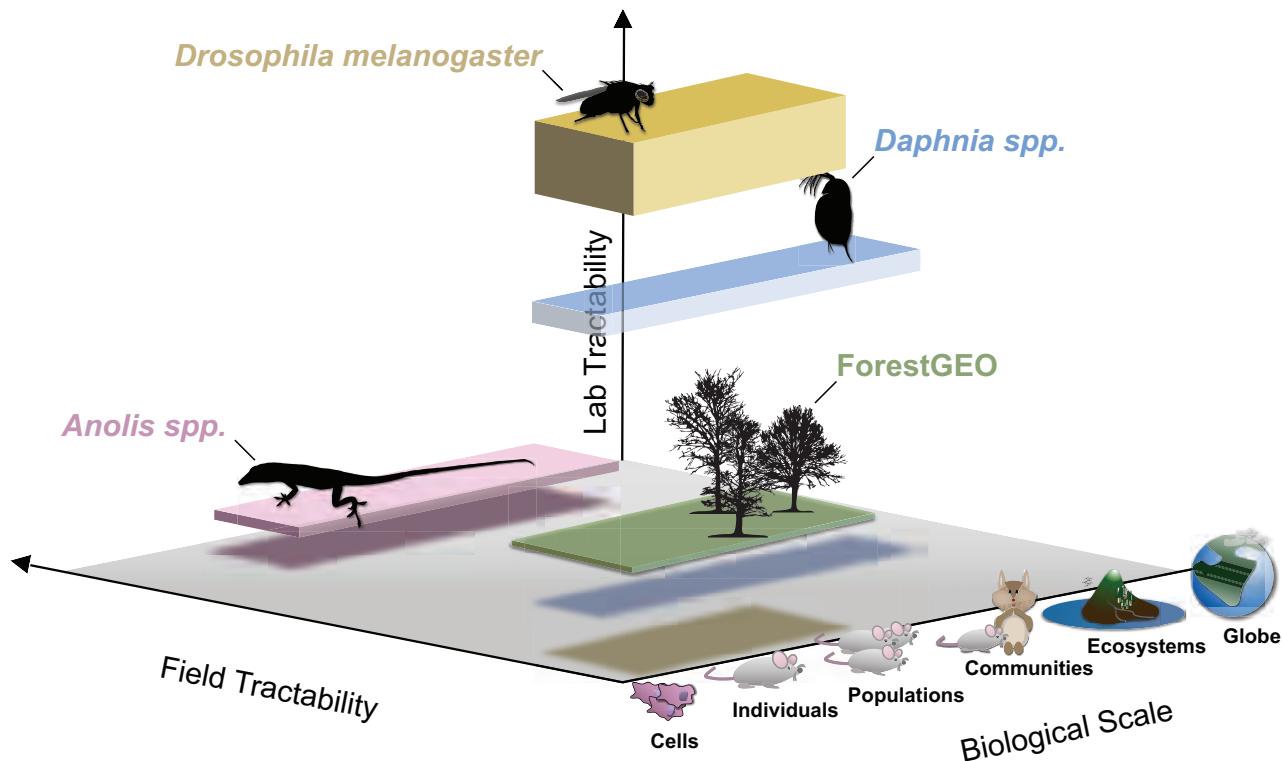


Figure 1: Model systems in ecology, evolution, and behavior (EEB) vary along a number of axes, including their tractability for field studies and for laboratory studies and the biological scales at which they are typically studied. This multidimensional view of space that model systems occupy harkens back to Hutchinson’s concept of the niche and his classic depiction of squirrels in an n -dimensional hypervolume (Hutchinson 1978). A model system’s position in this space influences the questions it is best suited to address. Further extending the Hutchinsonian metaphor, the realized space of a model system depends both on the biological features (i.e., fundamental aspects) and on the history of accumulated knowledge and techniques related to a particular system. In this figure, we represent approximately where a subset of common model systems in EEB fall along these three axes (see table S1). Organism silhouettes are from PhyloPic and 4vector. ForestGEO = Forest Global Earth Observatory.

Box 1: Skills and other attributes associated with developing new model systems, nurturing nascent model systems, and using existing systems in new ways

While people sometimes take the existence of model systems for granted, developing new model systems, nurturing nascent model systems, and using existing systems in new ways requires skills on the part of individual researchers and is facilitated by certain attributes of institutions and of the study system. We describe some particularly important attributes in this box.

Individual Attributes

Developing a new model system requires insight—what are the major gaps in our knowledge? What are major outstanding questions? What tools and methodologies can be leveraged to address those questions? What is the potential of a particular organism or system? And, just as importantly, what are its limitations? It also requires foresight—where is the field headed? Are there new technologies on the horizon that will open up major new research opportunities?

Developing a new model system also requires strong natural history skills, including excellent observational skills and record keeping. It requires an ability to tinker—having the curiosity, ingenuity, resourcefulness, and instincts to modify aspects of the environment or setup in a way that facilitates studies within a particular system. And unquestionably it requires an ability to persevere through setbacks—something that can be greatly facilitated by particular institutional structures, as we discuss more below.

Developing model systems also requires being a good collaborator, mentor, and communicator. Collaboration will promote studies of the same system by multiple researchers, which is required for building the depth of knowledge needed to achieve the “model system” designation. These collaborations are facilitated by a culture of openly sharing data, protocols, and other materials (Ankeny and Leonelli 2020; Matthews and Vosshall 2020) and by strong communication skills. A scientist who has an amazing vision but is unable to communicate that with others (including potential funders, collaborators, students, and others) will have limited impact. Networking skills are also useful, as they can help develop connections that allow for new lines of study in a particular system and that recruit more people to work on the system.

Institutional and Structural Attributes

Model systems are extensively studied from a variety of angles, yielding deep knowledge of that system. Thus, developing a model system is supported by having a diversity of researchers who work on the same system but approach it from different angles (or subdisciplines), using different techniques and approaches and with different perspectives. Crucially, EEB will only benefit from those diverse perspectives if our departments, field stations, meetings, and all of the other places where we do our work are inclusive spaces.

Funding is also a crucial component of developing new model systems. Building deep knowledge of a system requires many years of study by many people—something that can be achieved only with financial support. A major challenge in today’s funding climate is supporting work on the natural history of a system and funding that supports high-risk/high-reward projects.

A related factor is that there need to be structures in place that protect an individual researcher from the impacts of failures, such as job stability, supportive mentors, and other systems (e.g., evaluating candidates on the basis of a few publications of their choosing rather than their total number of publications). It is not clear whether there is a particular time in one’s career where it is “best” to develop a new model system; some researchers begin developing them relatively early in their careers, while others wait until they have already established themselves. As with so many things, a wide range of circumstances (not to mention serendipity) will play an important role in the timing.

Organism or Study System Attributes

Some organisms (or communities or ecosystems) are more readily established as model systems, based on factors such as ease of working with them in the field and/or laboratory, generation time, organism size, and population

Box 1 (Continued)

abundances. An organism that is small, abundant, reproduces quickly, and grows well in the field and the lab is more likely to become established as a model system than an organism that is large, rare, and difficult to grow. However, while there are challenges with organisms with more complex life histories (e.g., parasites that must pass through multiple hosts, organisms with biennial or multiannual life cycles), model systems that capture these diverse realities are essential for addressing fundamental questions in ecology and evolution. One possibility is to assemble longer-term data sets over time, with new members of a lab analyzing data collected by prior lab members and “paying it forward” by collecting additional data.

Using Existing Systems in Novel Ways

There is strong overlap between the skills needed to develop a model system and those needed to take an existing system and use it in a novel way, including insightfulness, a sense of where the field is heading, and good communication skills. Two additional attributes that are particularly important for using model systems in innovative ways are creativity and big-picture thinking, both of which enable a scientist to see beyond the scope of how a system has been used in the past. Without these, it is easy to remain within the confines of what has already been done rather than to use them as a foundation for a leap off in a new and exciting direction. Some useful questions to ask in the context of taking existing systems in new directions include: Are there modes of inquiry from other disciplines or modes of thought that could be newly applied to this system? How could our knowledge of an existing system change as a result of these new perspectives?

Finally, we note that serendipity can definitely play a role. Sometimes, model systems begin to be used in a new way because of a chance observation that occurs during a study of an entirely different question. However, these serendipitous occurrences will lead to new directions only if the attributes listed above are present. As Louis Pasteur put it: chance favors only the prepared mind.

Developing New Systems

As discussed elsewhere in this article, many traditional model systems were developed by people in positions of power (e.g., at traditionally powerful and wealthy institutions), and work on those systems is sometimes deemed important or worthy simply by virtue of being done in an established model system. We call on our community to use more of a bottom-up or community organizing approach as novel model systems are developed, getting buy-in from diverse members of our community.

Forest Global Earth Observatory (ForestGEO). At a smaller scale, mesocosms and microbial communities and ecosystems have emerged as model systems to study community ecology (Datta et al. 2016; Goldford et al. 2018; Fugère et al. 2020), species interactions (Mickalide and Kuehn 2019), ecosystem processes (de Jesús Astacio et al. 2020), and eco-evolutionary dynamics (Lawrence et al. 2012; Matthews et al. 2016). These types of systems allow us to incorporate ecological context and dynamics while still maintaining tractability (Sanchez et al. 2021). While these types of systems are not part of the traditional definition of a model system, they can be used to understand particular biomes and general principles at a global scale and allow us to avoid some of the biases that are associated with more traditional model systems (Bolker 1995; Alfred and Baldwin 2015). Our

definition of a model system is agnostic about the degree to which an organism (or community or ecosystem) is “representative.” Any one system will be representative of some aspects of ecology and evolutionary biology and unusual in others. Research in natural and experimental contexts provides insights into fundamental processes in EEB (Bartholomew 1966), as does work on organisms that are representative and those that are unusual; there can be as much to learn from a system that is an exception to a rule as from one that adheres to it.

Even though variation is a key focus of research in EEB, model systems are generally chosen and constructed in a way in which variation (or its drivers) are restricted or delimited. It is no accident that the ecosystems that have emerged as model systems are often delimited places, such

as islands or field plots; this isolation limits the contribution of “undesirable” variation or noise. Similarly, when the model system is an organism rather than an ecosystem, we tend to select organisms that we can standardize and isolate, such as by growing them in the laboratory (table S1; fig. 1). For this reason, model systems in EEB share some of the (biased) biological traits of model organisms in the biomolecular sciences that make them intrinsically tractable, such as ease of husbandry in laboratory conditions, fast generation times, and traits that are easily quantified (such as external color variation).

After defining what a model system is, it is worth considering what is not a model system. Systems that do not yet have the technology and knowledge base to allow for horizontal and vertical integration are not model systems. Sometimes, this is due to a lack of research on a particular system. Other times, this is because aspects of the system (e.g., life-history traits) pose challenges, especially given current institutional structures. While people have sometimes found creative solutions to working on such systems (e.g., with periodical cicadas; Yang 2004), current tenure review processes and models for funding graduate students can make it challenging to work on longer-lived organisms or on longer-term phenomena (box 1). Importantly, something that is a model system for one subarea of EEB is not necessarily a model system for all model areas (although certainly particular systems can be models for multiple areas). By our definition, systems can move from nonmodel to model status once a sufficient knowledge base has been developed and recognized by the community; systems cannot move from model to nonmodel status, although certain model systems may fall out of favor or stop being the subject of study because of other concerns (e.g., biosecurity; Wale and Duffy 2021).

As model systems become established, a positive feedback loop can kick in where research on the system makes additional research more likely (Matthews and Vosshall 2020). The development of standardized (and shared) knowledge about the system—protocols, natural history knowledge, techniques for quantifying variation, stock lines, mathematical models, and so on—makes these systems more tractable for additional research and facilitates the expansion of work on the system into new subdisciplines and new questions (box 1). Over time, there is greater vertical integration and more possibilities for horizontal comparison. The interconnection and integration between subdisciplines that arises as a result is a major strength of working with model systems.

At the same time, this expansion of research both through time and across subdisciplines can be particularly susceptible to the propagation of unsupported assumptions and erroneous inferences made early in the study of the system, potentially resulting in substantial bodies of work that rest on shaky foundations. For especially long-studied model

systems, some of these decades-old assumptions may be signs of the times, projections of entrenched sociocultural and political values onto the study system that no one thought to question at the time (Haraway 1989, 1991). Regardless of their source, these assumptions and inferences can become embedded into research on the model system and become challenging to escape, even as their impact is magnified throughout the field through horizontal and vertical integration. For example, *Anolis* lizards are a model system for studying convergent evolution and adaptive radiation and have been the subject of research in behavioral and evolutionary ecology for more than a century (Losos 2009). This research includes more than a hundred articles published on territorial behavior in *Anolis*. Through a comprehensive close reading to evaluate evidence for territoriality in these articles, Kamath and Losos (2017) revealed that territoriality was assumed rather than tested in the earliest research on anoles, and this early assumption became entrenched in subsequent work in this system, implicitly and explicitly shaping study design, data analysis, the interpretation of results, and publication. While similar assumptions have likely been made in many other taxa described as territorial, their origins and consequences were readily traceable in *Anolis* only because of the long history of research in this model system.

Long-studied model systems can be a compelling context in which to apply methods from the humanities and social sciences to understand scientific practice (e.g., Haraway 1989; Kohler 1994; Rader 2004; Milam 2010). Such work makes explicit the ever-present feedbacks between the questions we scientists ask and the identities, cultures, and socio-political contexts we bring to our work. This kind of cross-disciplinary inquiry into model systems can seed ideas for novel conceptual and empirical approaches to long-studied questions in EEB (Kamath and Losos 2018; Kamath and Wesner 2020). Equally, because the assumptions and inferences made early in the study of model systems can be deeply consequential, scientists working to establish new model systems would do well to consider the value of insights from cross-disciplinary inquiry for their work, including through formal collaborations with scholars in the social sciences and humanities who study the human dimensions of scientific practice. In this way, model systems can make room for disciplinary and methodological diversity in our study of the natural world.

Our Traditional Model Systems Reflect Historical Biases

Model systems not only allow us to answer scientific questions but also to play an important role in shaping the questions asked (Ankeny 2001; Leonelli 2007). This means that the history that shaped the establishment of our

Box 2: Model systems in the tropics

The establishment of model systems in the tropics was strongly influenced by sociopolitical context and ease of access to researchers from the United States. As a result of the Spanish-American war, the United States colonized not only Cuba but also Puerto Rico, Guam, and the Philippines. In 1904, the United States took formal control of the Panama Canal after actively supporting the separation of Panama from Colombia. Aided by the increased influence and control in the region, US institutions established different research sites in Central America and the Caribbean (e.g., Cinchona in Jamaica, the Harvard Botanical Garden in Cuba, and Barro Colorado Island in the Panama Canal region; Raby 2017).

After the Cuban revolution in 1959, Atkins Garden—at the time a main research center in the Neotropics—was forced to close its doors (Raby 2017). The National Science Foundation, together with the Organization of American States, sponsored three meetings to create a strategy to facilitate research of US scientists in tropical research (Stone 1988). The result of these meetings was the formation of the Organization for Tropical Studies (OTS), a consortium of universities and research centers in Latin America and the United States, with field stations in Costa Rica and South Africa (Rocha and Braker 2021).

OTS has offered the field course Fundamentals of Tropical Biology since 1961. The origin and popularization of many tropical model systems can be traced to research performed in OTS field stations and OTS field courses. Examples of classic model systems developed in OTS stations include *Heliconius* butterflies, *Piper* shrubs, army ants, interactions between leafcutter ants and associated microorganisms, and interactions between Zingiberales “banana-like plants” and rolled-leaf beetles (Rettenmeyer 1963; Gilbert 1972; Strong 1977; Marquis 1984). This history helps explain why studies in Costa Rica are highly overrepresented given its size (Stocks et al. 2008; Martin et al. 2012).

Many tropical model systems, including those still in use today, were developed by researchers from the Global North. While there is still a problem of underrepresentation of people from tropical countries as active participants in the science conducted there (Stocks et al. 2008), the efforts of OTS to promote inclusion of tropical scientists for more than 50 years has led to a growing number of researchers from the Global South working on these classical tropical systems (Chaves-Campos 2003; Mavárez et al. 2006; Pinto-Tomás et al. 2009; García-Robledo et al. 2016). Although many scientists in the Global South are playing central roles in research involving classic tropical model systems, men continue to outnumber women (Hill et al. 2010). In addition to the stereotypes and implicit biases that reduce participation of minorities in STEM, Latina scientists have to face the challenges associated with culturally ingrained concepts of masculinity (“machismo”; Bernal et al. 2019).

model systems has real effects on our science today and makes it important to consider the biases and historical contingencies associated with their establishment. Particular organisms become model systems not only because of their biology but also because of a variety of other factors, including the institutional structures that support them. Indeed, whole institutions have been created for the development, domestication, and standardization of traditional model systems and their associated protocols (Burian 1993; Clause 1993; Kohler 1994; Leonelli 2007).

Traditional model systems are generally highly constructed (e.g., genetic lines are carefully bred, standard breeding conditions carefully designed, and unwanted variation selected out). Thomas Hunt Morgan (1866–1945) and his group, for example, developed standardized protocols to grow *Drosophila* (minimizing variation in the expression of phenotypic traits) and made genetic lines enriched for differences in Mendelian inherited traits (Kohler 1994). Similarly, the reference strain of *Saccharomyces cerevisiae* that provided the foundation for early research in this system was generated via laboratory crosses

and selected because it was unusual in that it could be maintained as a haploid, facilitating the study of mutations (Liti 2015). These goals of control and technological development were closely linked to the increasing use of genetics for domestication, and the eugenic desires to control the genetic makeup of domesticated animals, crops, and humans for the “betterment of society” (Bowman and Rebolledo-Gómez 2020).

In addition, because science was dominated by Western scientific institutions, traditional model systems were often chosen because they were easy to access and amenable to study by scientists working at those institutions. *Drosophila*, for example, was chosen as a model system because its phenology made work on it convenient given the academic calendar in the northern United States; they were most abundant in fruit orchards early in the fall, and students could easily breed them indoors during the winter (Kohler 1994). The common house mouse (*Mus musculus*) was common in Europe and industrialized cities in the United States, and in addition mice were bred by mice fanciers for their rare coat colors and odd behaviors; thus, lines of mice bred

Box 3: How to assess whether there are gaps in existing model systems

We propose that the general approach used by Wale and Duffy (2021) can provide a framework for evaluating whether existing model systems in use in a given subdiscipline are sufficient or whether the subdiscipline would benefit from additional systems.

Evaluating Currently Used Systems

Step 1. Identify the key processes and phenomena of interest to a subdiscipline. For example, existing theory on the ecology and evolution of infectious diseases points to three processes—transmission, disease, and recovery—as fundamental. Making these key processes and phenomena explicit also allows for researchers to add or modify them, which can be an important way in which research in a subdiscipline progresses.

Step 2. Review the current model systems that are in use in that subdiscipline. This review should focus on assessing whether each individual study of a given system explores the fundamental processes identified in step 1. While carrying out the review, it is likely that additional important features and differences will emerge (e.g., related to the scale at which particular processes are studied in particular systems).

Step 3. Using the results from step 2, evaluate whether the systems currently in use in that subdiscipline are capturing a wide range of parameter space for the processes of interest. Can the systems currently in use illuminate core themes and processes for that subdiscipline (Jenner and Wills 2007)?

Step 4. What are the underlying assumptions about existing model systems? Have those assumptions been tested?

Step 5. Consider whether, in addition to the key processes identified, there are other notable gaps in the model systems currently in use. One that is likely to be true in many subdisciplines is that existing model systems might come from a relatively restricted geographical area or may represent only certain life-history traits (table S1). We propose considering how broadly you can apply knowledge using current systems. Does it tell you about only a certain type of organism or ones that live in certain locales? How well is the parameter space in figure 1 covered? Similar to what is often done with mathematical models, it is important to be explicit about what our model systems represent and, even more importantly, what they do not represent.

Steps to Take If (or, More Likely, When) Gaps Are Identified

Here are some questions to ask when trying to identify systems that might be developed and used to fill existing gaps.

1. Are there model systems in use in other areas of ecology, evolutionary biology, and behavior (or, if not, other areas of biology) that can help fill those gaps?
2. Are there nascent study systems that are promising—for example, ones that have been studied in nature for a long time but that would benefit from development of novel molecular tools?
3. Can the model systems under consideration be manipulated and studied on the timescales of a PhD program or while an assistant professor? If not, how have others who work on organisms or processes with longer timescales approached those questions?
4. Can museum collections be of use, including to extend temporal and/or spatial scales? Consider, however, the likelihood of biased representation within museum collections (Loiselle et al. 2007; Wehi et al. 2012; Gower et al. 2019; Thompson and Birkhead 2020).
5. What sources of information might exist outside those typically considered by Western scientists? Are there other historical records (e.g., phenological data collected by community scientists or existing photo or video collections) that can be used to address the question? What do local communities already know about the system? What work has been done on the topic by non-Western scholars (including work published in languages other than English)?

Checkpoint. When considering the development of a new potential model system, it is essential to ask whether it will be done in a way that increases or decreases inequity. Unfortunately, there is a long history of extractive practices that reinforce colonialism and imperialism (DuBay et al. 2020; Gewin 2021), of research that “discovers”

Box 3 (Continued)

things that were already well known in local communities (e.g., Cañizares-Esguerra 2019), and of research that ignores the contribution of non-Western scientists (e.g., Malik et al. 2018).

Researchers should also consider whether their work would benefit from establishing multiple taxa at the same time (depending on the study topic, these might be chosen because they are closely related or, alternatively, because they encompass phylogenetic breadth).

for clear phenotypic characteristics were commercially available. At the time when the mouse was becoming a model for the study of genetics, there was a good market for “mouse fancy” in New England that allowed Castle and Little to start their genetic studies in mice with lines from a farm in Massachusetts (Rader 2004).

At the same time as these traditional biomolecular model systems became well developed, the establishment of modern academic ecology was accompanied by the extensive study and establishment of particular ecological ecosystems as models (e.g., work by Henry Cowles [1869–1939] on succession in the Indiana Dunes, research by Raymond Lindeman [1915–1942] in Cedar Bog Lake [part of what is now Cedar Creek Ecosystem Reserve], and work by G. Evelyn Hutchinson [1903–1991] on Linsley Pond; Golley 1993). Over time, there was a growing awareness of the geographical biases in where ecological research was being performed and a desire to do more systematic research in the tropics (Richards 1963). One consequence of this was that the number of field stations rapidly increased (Tydecks et al. 2016), but in a way that was uneven and that still reflected ease of access by researchers from the United States. This contributed to the substantial overrepresentation (given their size) of research done in Panama and Costa Rica (Stocks et al. 2008; Martin et al. 2012). The uneven establishment of field stations in the tropics was strongly impacted by the geopolitical context (box 2). Despite efforts to expand the geographic range of research in EEB, most of the research published in the major ecological journals is still based on sites in Europe and in the United States (Martin et al. 2012).

It is clear that there are strong historical and systemic biases impacting the classic model systems in EEB, as well as clear gaps in our existing model systems (box 3; table S1). Recently, there has been a push to expand and diversify the use of models by including more female animals in biomedical studies (Shansky 2019), including more phylogenetic diversity around well-studied model organisms and traits of interest (Jenner and Wills 2007), and adding more ecological complexity in our systems (Rillig and Antonovics 2019; Sanchez et al. 2021). Filling the gaps in existing model systems will also require a concerted effort by researchers and funding agencies to invest in the re-

sources (including establishing strain databases, molecular toolkits, and computational software) and studies of natural history that facilitate research using emerging model systems (Matthews and Vosshall 2020; box 1).

Overall, model systems emerge from the community as a result of countless decisions made by individual scientists (including early-career scientists; box 4), with a strong influence of our institutional cultures. Increasing buy-in from the community is often beneficial for the model system (and the scientists whose careers are tied to these models), but these communities can also serve as gatekeepers. Therefore, in addition to focusing on diversity of our model systems, we must focus on diversity and inclusion in our discipline. Indeed, when new model systems are built with intention, this can be a mechanism for increasing diversity and inclusion in EEB (box 4).

EEB Needs Diverse Scientists

STEM disciplines were designed for one particular type of person—white men who are cisgendered, heterosexual, not disabled, and from relatively affluent backgrounds (McGee 2020). EEB as a discipline was also designed for this type of person and, like STEM, more broadly has been—and still is—in hospitable to people who do not fit that mold (Valantine et al. 2016; Graves 2019; Kaishian and Djoulakian 2020; McGee 2020; Montgomery 2020a; Wanelik et al. 2020). This is especially true for scientists who hold multiple minoritized identities (Ireland et al. 2018). It is important to note that despite these barriers, scientists from underrepresented groups have long made contributions to EEB (Bronstein and Bolnick 2018; Mackay et al. 2019; Jaffe et al. 2020; Lee 2020).

While we would benefit from more comprehensive data, it is clear that the demographics of EEB still reflects these origins and this exclusion. To give some examples: according to the US National Science Foundation’s Survey of Earned Doctorates, 387 US citizens and permanent residents earned PhDs in ecology in 2019, and 192 earned PhDs in evolutionary biology (NCSES 2019). Of those who earned PhDs in ecology, 322 were White and not Hispanic or Latino (~83% of the total); the comparable number for evolutionary biology was 146 (~76% of the total). Only

Box 4: Additional considerations for early-career researchers

As early-career scientists establish their careers, they must make decisions about what systems to study. Making these decisions often involves considerations beyond just the scientific questions they are interested in tackling. Will they have access to the necessary resources? Will the field be welcoming? Will they be able to carve out a niche of their own?

A key challenge for early-career scientists is how to differentiate from previous mentors and other established groups. How does a seedling lab carve out its space in a crowded forest? Even in cases where the community working on a particular model system is welcoming and eager to share resources, early-career researchers face challenges in establishing their independent groups. The reality is likely to be that seedling labs will have relatively few resources (in terms of both people and funding), while the research forest might have some large trees that cast a very long shadow. And, unfortunately, the research environment for particular model systems is not always welcoming, especially for early-career scientists who do not fit the traditional “model scientist” mold. A key aspect of working on model systems is the community associated with it, which provides a variety of perspectives (e.g., from different subfields of EEB), can share protocols and help someone learn new techniques, and can help an early-career researcher negotiate a distinct intellectual niche within that research community. Whether scientists from underrepresented groups are less (or more) likely to work on model systems would be an interesting topic for further investigation.

As a result, in some cases early-career scientists will decide that the best path forward is to establish a new model system or to take an existing model system and use it in a very different way. This has the advantage of avoiding competition. Establishing a new system (or using an existing one in a very different way) also can mean that work on a particular system (or in a particular subdiscipline) is done by diverse scientists from the start, with the potential to establish a healthy, equitable, and inclusive culture right from the beginning.

However, as always in ecology and evolution and in life, there are trade-offs. While there are advantages to establishing new systems, there are also important drawbacks. First, there is a larger-than-average chance of failure when trying to do something completely new; deciding whether to take on this risk at a particularly vulnerable career stage will require careful thought. Second, establishing new systems will require funding, including for natural history work and for work that is high-risk/high-reward, neither of which are well supported in current funding climates (as also mentioned in box 1). Third, moving into a new model system from an established model system may lead to a loss of research connections and community, including potentially an impact on the rate at which articles are cited.

eight (1.4%) PhD recipients in ecology and evolutionary biology in 2019 were Black, and only one (0.2%) was Native American (NCSSES 2019). In New Zealand, Māori and Pasifika are severely underrepresented at the faculty level at universities and crown-research institutes, with little progress over a decade (McAllister et al. 2020). Survey responses from attendees at the Evolution 2019 meeting indicated that the representation of women drops with career stage, as does representation of LGBTQ+ scientists (Rushworth et al. 2021); consistent with this, women scientists tend to have shorter publishing careers (Huang et al. 2020). An analysis of top-publishing authors in ecology, evolution, and conservation found that only 11% are women and that 10 countries from the Global North (inclusive of Australia) account for 86% of top-publishing authors (Maas et al. 2021). There is also strong geographic bias in the composition of editorial boards in ecology, evolutionary biology, and closely related fields; an analysis of the editorial boards of 20 leading conservation biology

journals revealed that they had few or no editors from regions with the most biodiversity (Campos-Arceiz et al. 2018). Unfortunately, it is clear that our field is still far from being diverse, equitable, and inclusive.

The lack of diversity in EEB holds back our science (Ireland et al. 2018; Duc Bo Massey et al. 2021). People with different backgrounds and lived experiences will approach science differently, asking different questions and pursuing different lines of research (Keller 1982; Stewart and Valian 2018; Duc Bo Massey et al. 2021). The science we do—the questions we ask and how we pursue answers—is influenced by our identities and by the social and political context in which we were raised (Keller 1982; Harding 1986; Wall Kimmerer 2013; Duc Bo Massey et al. 2021). Because gatekeepers often share many of the identities and backgrounds with the traditional “model scientist,” many scientists who did not fit that mold were told that the questions they asked were “not science” (Keller 1982; Haraway 1989; Wall Kimmerer 2013), and surely many more who were told

this were driven away from science. This is a problem from a justice perspective, and it also means that science suffers. Students from underrepresented groups are more innovative than majority students, though unfortunately their innovations and contributions tend not to be recognized and appreciated (Hofstra et al. 2020). Moreover, for teams working together on a project, diverse groups outperform homogeneous ones (Hong and Page 2004)—a result that parallels findings in nonhuman communities (Tilman et al. 2001). Model systems research will benefit greatly from a more diverse community of researchers.

More importantly, the lack of diversity in EEB (and STEM more broadly) is a moral and ethical issue. While there are clear arguments that science benefits from diversity (as discussed above), scientists from underrepresented groups should have the same opportunities to do science, and these opportunities should not rest on appeals to exceptionalism or benefits to science. Everyone should have an opportunity to do science.

Increasing Diversity, Equity, and Inclusion in EEB Will Require Acting at Many Levels, Including Making Structural and Institutional Changes

If there is one loud and clear message from the research literature on workplace diversity, it is that multiple, interacting, nested levels of context matter. (Bond and Haynes 2014)

Increasing diversity in EEB and creating a discipline that is inclusive of people of all backgrounds and identities requires a focus on institutional structures and gatekeepers (McGee 2020). Many efforts to increase diversity in STEM disciplines focus on individual students, especially on preparing these students (which often translates into attempts to “fix” or assimilate students from underrepresented groups; Bowman and Rebolledo-Gómez 2020; Halsey et al. 2020; McGee 2020; Schell et al. 2020). Alternatively, conversations focus on the changes that will come as more diverse early-career scholars progress through the academic ranks, ignoring that this is not a simple issue of demography (Holman et al. 2018). Instead of viewing the lack of diversity through the problematic “pipeline” metaphor (Cannady et al. 2014; McGee 2020), we must focus on changing structures, including focusing on how racism (and other isms) within a department and institution underlie the lack of diversity (McGee 2020). If organisms that we study fail to grow or thrive in an environment, we consider what aspects of the environment might be causing that outcome (Montgomery 2020a, 2020b); it is essential that we do the same with marginalized and minoritized scientists and that we work to change our institutions (including our departments and scientific societies) so that they are inclusive and

enable scientists who are outside the traditional “model scientist” mold to thrive (McGee 2020; Montgomery 2020a, 2020b).

One major challenge in EEB relates to who has access to research opportunities; at present, such opportunities are often inaccessible to individuals who come from socioeconomically disadvantaged backgrounds. A lack of accessibility to field courses and fieldwork can prevent people from entering the field (Beltran et al. 2020; McGill et al. 2021). Moreover, positions where early-career scientists, including field and laboratory technicians, are expected to work pro bono (or even pay for the experience) excludes research participation by individuals unable to self-fund or work for free, which disproportionately cuts off research opportunities for individuals from underrepresented groups (Fournier and Bond 2015; Emery et al. 2019). Additional challenges include working toward developing an understanding (and respect) for the social, cultural, and environmental experiences shared among individuals belonging to underrepresented groups. Doing so will increase the absent sense of “belonging” for these individuals within EEB and academia and nurture a field wherein one does not feel they need to conform to the cultural norms instituted by gatekeepers to ensure successful careers (Duc Bo Massey et al. 2021; McGill et al. 2021).

Social science research demonstrates that organizational-level policies strongly influence the degree to which minoritized groups are fully integrated into that organization and points to changes that can be implemented to increase inclusion (Bond and Haynes 2014). These changes include clearly communicating that behaviors that discriminate against individuals from certain groups will not be tolerated and clearly indicating that the organization views diversity as an asset that is important to the (shared) mission of the organization and its employees (Bond and Haynes 2014). Moreover, changes need to occur at multiple levels—a person’s trajectory in science and the environment they experience are influenced by factors at multiple levels (Bond and Haynes 2014; Valentine et al. 2016; Zea and Bowleg 2016). For example, scientists from the Global South face major barriers even as immigrants in Europe, Canada, and the United States. Immigrant scientists and international students from nonprivileged backgrounds start their careers abroad at economic disadvantage, as a substantial portion of their income must be invested in fees associated with immigration. In addition to the influences of biased gatekeepers and departmental culture, institutional and federal funding structures make it more expensive for departments to support these students and further restricts access to key fellowships. An important additional consideration in EEB relates to field safety. Certain individuals are at greater risk of harm and conflict when carrying out fieldwork, and faculty, departments, and institutions must help people in their

labs evaluate these risks and consider strategies that can help mitigate them (Demery and Pipkin 2021).

Several recent articles highlight specific changes that can be made to promote diversity and inclusion in academia, STEM, and EEB. Some of these are aimed at people in majority groups—and especially at White faculty (Sensoy and DiAngelo 2017; Platt 2020; Schell et al. 2020; Stevens et al. 2021)—while others are aimed at scientists from underrepresented groups (Halsey et al. 2020; Tseng et al. 2020). One common theme is the importance of welcoming scholars from underrepresented groups to bring their authentic selves to their research, rather than expecting them to assimilate to majority cultural norms; as Schell et al. (2020) note, we appreciate and recognize the value of diversity in the ecosystems we study, yet we expect homogeneity and assimilation of those carrying out the work. For EEB to be truly inclusive—and for our science to benefit from diversity—marginalized voices need to be heard, centered, and amplified.

Diverse Scientists Will Yield Diverse Model Systems and Diverse Perspectives, Improving Our Understanding of Ecology and Evolutionary Biology

Our understanding of ecology and evolutionary biology is the product of thousands of individual decisions regarding what questions to ask and which systems to study. When those decisions are made by relatively homogenous groups and when our work focuses on relatively few taxa, the conclusions we draw will be limited, and our understanding constrained. If we wish to uncover general phenomena and processes in ecology and evolutionary biology, we must support and nurture work on many different model systems, and we must invite and welcome contributions from scientists of all backgrounds and identities. Diverse model systems and diverse scientists will provide diverse perspectives, which in turn will allow us to understand endless forms most beautiful.

Acknowledgments

We thank Aisha Burton for permission to use her tweet as the epigraph. We also thank two anonymous reviewers for their feedback on the manuscript and Robert Arlinghaus, Jeremy Fox, Sarah Knutie, Anna-Liisa Laine, Jonathan Losos, David Lowry, Levi Morran, Amy Pedersen, Ken Olsen, Tamal Roy, and Trisha Wittkopp for their feedback on table S1. M.A.D. acknowledges support from the Moore Foundation (GBMF9202; <https://doi.org/10.37807/GBMF9202>). C.G.-R. acknowledges support from the National Science Foundation (Dimensions of Biodiversity grant 1737778).

Statement of Authorship

All authors contributed to the framing, writing, and editing of this article.

Literature Cited

Alfred, J., and I. T. Baldwin. 2015. New opportunities at the wild frontier. *eLife* 4:e06956.

Ankeny, R. A. 2001. The natural history of *Caenorhabditis elegans* research. *Nature Reviews Genetics* 2:474–479.

Ankeny, R. A., and S. Leonelli. 2011. What's so special about model organisms? *Studies in History and Philosophy of Science A* 42:313–323.

Ankeny, R. A., and S. Leonelli. 2020. *Model organisms*. Cambridge University Press, Cambridge.

Bartholomew, G. A. 1966. *Interaction of physiology and behavior under natural conditions. The Galapagos*. University of California Press, Berkeley, CA.

Beltran, R. S., E. Marnocha, A. Race, D. A. Croll, G. H. Dayton, and E. S. Zavaleta. 2020. Field courses narrow demographic achievement gaps in ecology and evolutionary biology. *Ecology and Evolution* 10:5184–5196.

Bernal, X. E., B. Rojas, M. A. Pinto-E, Á. M. Mendoza-Henao, A. Herrera-Montes, M. I. Herrera-Montes, A. Del Pilar Cáceres Franco, et al. 2019. Empowering Latina scientists. *Science* 363:825–826.

Bolker, J. A. 1995. Model systems in developmental biology. *BioEssays* 17:451–455.

Bond, M. A., and M. C. Haynes. 2014. Workplace diversity: a social-ecological framework and policy implications. *Social Issues and Policy Review* 8:167–201.

Bowman, M., and M. Rebollo-Gómez. 2020. Uprooting narratives: legacies of colonialism in the neoliberal university. *Hypatia* 35:18–40.

Bronstein, J. L., and D. I. Bolnick. 2018. “Her joyous enthusiasm for her life-work . . .”: early women authors in *The American Naturalist*. *American Naturalist* 192:655–663.

Burian, R. M. 1993. How the choice of experimental organism matters: epistemological reflections on an aspect of biological practice. *Journal of the History of Biology* 26:351–367.

Campos-Arceiz, A., R. B. Primack, A. J. Miller-Rushing, and M. Maron. 2018. Striking underrepresentation of biodiversity-rich regions among editors of conservation journals. *Biological Conservation* 220:330–333.

Cañizares-Esguerra, J. 2019. Screw Humboldt. Medium. <https://jorgecanizaresesguerra.medium.com/screw-humboldt-def1320213f5>.

Cannady, M. A., E. Greenwald, and K. N. Harris. 2014. Problematizing the STEM pipeline metaphor: is the STEM pipeline metaphor serving our students and the STEM workforce? *Science Education* 98:443–460.

Chaves-Campos, J. 2003. Localization of army-ant swarms by ant-following birds on the Caribbean slope of Costa Rica: following the vocalization of antbirds to find the swarms. *Ornitología Neotropical* 14:289–294.

Clause, B. T. 1993. The Wistar Rat as a right choice: establishing mammalian standards and the ideal of a standardized mammal. *Journal of the History of Biology* 26:329–349.

Datta, M. S., E. Sliwerska, J. Gore, M. F. Polz, and O. X. Cordero. 2016. Microbial interactions lead to rapid micro-scale successions on model marine particles. *Nature Communications* 7:11965.

de Jesús Astacio, L. M., K. H. Prabhakara, Z. Li, H. Mickalide, and S. Kuehn. 2020. Closed microbial communities self-organize to

persistently cycle carbon. *bioRxiv*, <https://doi.org/10.1101/2020.05.28.121848>.

Demery, A.-J. C., and M. A. Pipkin. 2021. Safe fieldwork strategies for at-risk individuals, their supervisors and institutions. *Nature Ecology and Evolution* 5:5–9.

Dietrich, M. R., R. A. Ankeny, N. Crowe, S. Green, and S. Leonelli. 2020. How to choose your research organism. *Studies in History and Philosophy of Biological and Biomedical Sciences* 80:101227.

DuBay, S., D. H. Palmer, and N. Piland. 2020. Global inequity in scientific names and who they honor. *bioRxiv*, <https://doi.org/10.1101/2020.08.12.254110>.

Duc Bo Massey, M., S. Arif, C. Albury, and V. A. Cluney. 2021. Ecology and evolutionary biology must elevate BIPOC scholars. *Ecology Letters* 24:913–919. <https://doi.org/10.1111/ele.13716>.

Emery, N., A. Hund, R. Burks, M. Duffy, C. Scoffoni, and A. Swei. 2019. Students as ecologists: strategies for successful mentorship of undergraduate researchers. *Ecology and Evolution* 9:4316–4326.

Fournier, A. M. V., and A. L. Bond. 2015. Volunteer field technicians are bad for wildlife ecology. *Wildlife Society Bulletin* 39:819–821.

Fugère, V., M.-P. Hébert, N. B. da Costa, C. C. Y. Xu, R. D. H. Barrett, B. E. Beisner, G. Bell, et al. 2020. Community rescue in experimental phytoplankton communities facing severe herbicide pollution. *Nature Ecology and Evolution* 4:578–588.

García-Robledo, C., and C. S. Baer. 2021. Demographic attritions, elevational refugia, and the resilience of insect populations to projected global warming. *American Naturalist* 198:113–127.

García-Robledo, C., E. K. Kuprewicz, C. L. Staines, T. L. Erwin, and W. J. Kress. 2016. Limited tolerance by insects to high temperatures across tropical elevational gradients and the implications of global warming for extinction. *Proceedings of the National Academy of Sciences of the USA* 113:680–685.

Gewin, V. 2021. Respect and representation: Indigenous scientists seek inclusion for their knowledge and for themselves. *Nature* 589:315–317. <https://media.nature.com/original/magazine-assets/d41586-021-00022-1/d41586-021-00022-1.pdf>.

Gilbert, L. E. 1972. Pollen feeding and reproductive biology of *Heliconius* butterflies. *Proceedings of the National Academy of Sciences of the USA* 69:1403–1407.

Goldford, J. E., N. Lu, D. Bajić, S. Estrela, M. Tikhonov, A. Sanchez-Gorostiaga, D. Segrè, et al. 2018. Emergent simplicity in microbial community assembly. *Science* 361:469–474.

Golley, F. B. 1993. A history of the ecosystem concept in ecology: more than the sum of the parts. Yale University Press, New Haven, CT.

Gordon, S. P., E. Burdfield-Steel, J. Kirvesoja, and J. Mappes. 2021. Safety in numbers: how color morph frequency affects predation risk in an aposematic moth. *American Naturalist* 198:128–141.

Gower, G., L. E. Fenderson, A. T. Salis, K. M. Helgen, A. L. van Loenen, H. Heiniger, E. Hofman-Kamińska, et al. 2019. Widespread male sex bias in mammal fossil and museum collections. *Proceedings of the National Academy of Sciences of the USA* 116:19019–19024.

Grant, N. A., R. Maddamsetti, and R. E. Lenski. 2021. Maintenance of metabolic plasticity despite relaxed selection in a long-term evolution experiment with *Escherichia coli*. *American Naturalist* 198:93–112.

Graves, J. L. 2019. African Americans in evolutionary science: where we have been, and what's next. *Evolution: Education and Outreach* 12:18.

Green, D. A., II. 2021. Monarch butterfly migration as an integrative model of complex trait evolution. *American Naturalist* 198:142–157.

Green, S., M. R. Dietrich, S. Leonelli, and R. A. Ankeny. 2018. “Extreme” organisms and the problem of generalization: interpreting the Krogh principle. *History and Philosophy of the Life Sciences* 40:65.

Halsey, S. J., L. R. Strickland, M. Scott-Richardson, T. Perrin-Stowe, and L. Massenburg. 2020. Elevate, don’t assimilate, to revolutionize the experience of scientists who are Black, Indigenous and people of colour. *Nature Ecology and Evolution* 4:1291–1293.

Haraway, D. J. 1989. *Primate visions: gender, race, and nature in the world of modern science*. Psychology Press, London.

———. 1991. *Simians, cyborgs, and women: the reinvention of nature*. Routledge, New York.

Harding, S. G. 1986. *The science question in feminism*. Cornell University Press, Ithaca, NY.

Hill, C., C. Corbett, and A. St Rose. 2010. Why so few? women in science, technology, engineering, and mathematics. *American Association of University Women*, Washington, DC.

Hofstra, B., V. V. Kulkarni, S. Munoz-Najar Galvez, B. He, D. Jurafsky, and D. A. McFarland. 2020. The diversity-innovation paradox in science. *Proceedings of the National Academy of Sciences of the USA* 117:9284–9291.

Holman, L., D. Stuart-Fox, and C. E. Hauser. 2018. The gender gap in science: how long until women are equally represented? *PLoS Biology* 16:e2004956.

Hong, L., and S. E. Page. 2004. Groups of diverse problem solvers can outperform groups of high-ability problem solvers. *Proceedings of the National Academy of Sciences of the USA* 101:16385–16389.

Huang, J., A. J. Gates, R. Sinatra, and A.-L. Barabási. 2020. Historical comparison of gender inequality in scientific careers across countries and disciplines. *Proceedings of the National Academy of Sciences of the USA* 117:4609–4616.

Hutchinson, G. E. 1978. *An introduction to population ecology*. Yale University Press, New Haven, CT.

Ireland, D. T., K. E. Freeman, C. E. Winston-Proctor, K. D. DeLaine, S. McDonald Lowe, and K. M. Woodson. 2018. (Un)hidden figures: a synthesis of research examining the intersectional experiences of Black women and girls in STEM education. *Review of Research in Education* 42:226–254.

Jaffe, K., J. C. Correa, and Z. Tang-Martínez. 2020. Ethology and animal behaviour in Latin America. *Animal Behaviour* 164:281–291.

Jenner, R. A., and M. A. Wills. 2007. The choice of model organisms in evo-devo. *Nature Reviews Genetics* 8:311–319.

Kaishian, P., and H. Djoulakian. 2020. The science underground. *Catalyst: Feminism, Theory, Technoscience* 6:1–26. <https://doi.org/10.28968/cftt.v6i2.33523>.

Kamath, A., and J. Losos. 2017. The erratic and contingent progression of research on territoriality: a case study. *Behavioral Ecology and Sociobiology* 71:89.

———. 2018. Reconsidering territoriality is necessary for understanding *Anolis* mating systems. *Behavioral Ecology and Sociobiology* 72:106.

Kamath, A., and A. B. Wesner. 2020. Animal territoriality, property and access: a collaborative exchange between animal behaviour and the social sciences. *Animal Behaviour* 164:233–239.

Katz, P. S. 2016. “Model organisms” in the light of evolution. *Current Biology* 26:R649–R650.

Keller, E. F. 1982. Feminism and science. *Signs: Journal of Women in Culture and Society* 7:589–602.

Kohler, R. E. 1994. *Lords of the fly: Drosophila genetics and the experimental life*. University of Chicago Press, Chicago.

Kokko, H. 2020. When synchrony makes the best of both worlds even better: how well do we really understand facultative sex? *American Naturalist* 195:380–392.

Krogh, A. 1929. The progress of physiology. *Science* 70:200–204.

Lawrence, D., F. Fiegna, V. Behrends, J. G. Bundy, A. B. Phillimore, T. Bell, and T. G. Barraclough. 2012. Species interactions alter evolutionary responses to a novel environment. *PLoS Biology* 10: e1001330.

Lee, D. N. 2020. Diversity and inclusion activism in animal behaviour and the ABS: a historical view from the U.S.A. *Animal Behaviour* 164:273–280.

Leonelli, S. 2007. Growing weed, producing knowledge: an epistemic history of *Arabidopsis thaliana*. *History and Philosophy of the Life Sciences* 29:193–223.

Leonelli, S., and R. A. Ankeny. 2013. What makes a model organism? *Endeavour* 37:209–212.

Lewontin, R. C. 2001. The triple helix: gene, organism, and environment. Harvard University Press, Cambridge, MA.

Liti, G. 2015. The fascinating and secret wild life of the budding yeast *S. cerevisiae*. *eLife* 4:e05835.

Loiselle, B. A., P. M. Jørgensen, T. Consiglio, I. Jiménez, J. G. Blake, L. G. Lohmann, and O. M. Montiel. 2007. Predicting species distributions from herbarium collections: does climate bias in collection sampling influence model outcomes? *Journal of Biogeography* 35:105–116.

Losos, J. B. 2009. Lizards in an evolutionary tree: ecology and adaptive radiation of anoles. University of California Press, Berkeley.

Maas, B., R. J. Pakeman, L. Godet, L. Smith, V. Devictor, and R. Primack. 2021. Women and Global South strikingly underrepresented among top-publishing ecologists. *Conservation Letters*, <https://doi.org/10.1111/conl.12797>.

Mackay, A. W., D. Adger, A. L. Bond, S. Giles, and E. Ochu. 2019. Straight-washing ecological legacies. *Nature Ecology and Evolution* 3:1611.

Malik, A. H., J. M. Ziermann, and R. Diogo. 2018. An untold story in biology: the historical continuity of evolutionary ideas of Muslim scholars from the 8th century to Darwin's time. *Journal of Biological Education* 52:3–17.

Marquis, R. J. 1984. Leaf herbivores decrease fitness of a tropical plant. *Science* 226:537–539.

Martin, L. J., B. Blossey, and E. Ellis. 2012. Mapping where ecologists work: biases in the global distribution of terrestrial ecological observations. *Frontiers in Ecology and the Environment* 10:195–201.

Matthews, B., T. Aebischer, K. E. Sullam, B. Lundsgaard-Hansen, and O. Seehausen. 2016. Experimental evidence of an eco-evolutionary feedback during adaptive divergence. *Current Biology* 26:483–489.

Matthews, B. J., and L. B. Vosshall. 2020. How to turn an organism into a model organism in 10 “easy” steps. *The Journal of Experimental Biology* 223:jeb218198.

Mavárez, J., C. A. Salazar, E. Bermingham, C. Salcedo, C. D. Jiggins, and M. Linares. 2006. Speciation by hybridization in *Heliconius* butterflies. *Nature* 441:868–871.

McAllister, T. G., S. Naepi, E. Wilson, D. Hikuroa, and L. A. Walker. 2020. Under-represented and overlooked: Māori and Pasifika scientists in Aotearoa New Zealand's universities and crown-research institutes. *Journal of the Royal Society of New Zealand*, <https://doi.org/10.1080/03036758.2020.1796103>.

McGee, E. O. 2020. Interrogating structural racism in STEM higher education. *Educational Researcher* 49:633–644.

McGill, B. M., M. J. Foster, A. N. Pruitt, S. G. Thomas, E. R. Arsenault, J. Hanschu, K. Wahwahsuck, et al. 2021. You are welcome here: a practical guide to diversity, equity, and inclusion for undergraduates embarking on an ecological research experience. *Ecology and Evolution* 11:3636–3645. <https://doi.org/10.1002/ece3.7321>.

Mickalide, H., and S. Kuehn. 2019. Higher-order interaction between species inhibits bacterial invasion of a phototrophic-predator microbial community. *Cell Systems* 9:521–533.

Milam, E. L. 2010. Looking for a few good males: female choice in evolutionary biology. Johns Hopkins University Press, Baltimore.

Montgomery, B. L. 2020a. Lessons from microbes: what can we learn about equity from unculturable bacteria? *mSphere* 5: e01046-20.

———. 2020b. Planting equity: using what we know to cultivate growth as a plant biology community. *Plant Cell* 32:3372–3375.

NCSES (National Center for Science and Engineering Statistics). 2019. Doctorate recipients from U.S. universities: 2019. No. NSF 21-308.

Penczykowski, R. M., and R. D. Sieg. 2021. *Plantago* spp. as models for studying the ecology and evolution of species interactions across environmental gradients. *American Naturalist* 198:158–176.

Pinto-Tomás, A. A., M. A. Anderson, G. Suen, D. M. Stevenson, F. S. T. Chu, W. W. Cleland, P. J. Weimer, et al. 2009. Symbiotic nitrogen fixation in the fungus gardens of leaf-cutter ants. *Science* 326:1120–1123.

Platt, M. O. 2020. We exist. We are your peers. *Nature Reviews Materials* 5:783–784.

Raby, M. 2017. American tropics: the Caribbean roots of biodiversity science. University of North Carolina Press, Chapel Hill.

Rader, K. 2004. Making mice: standardizing animals for American biomedical research, 1900–1955. Princeton University Press, Princeton, NJ.

Rettenmeyer, C. W. 1963. Behavioral studies of army ants. University of Kansas Science Bulletin 44:281–465.

Richards, P. W. 1963. What the tropics can contribute to ecology. *Journal of Ecology* 51:231–241.

Rillig, M. C., and J. Antonovics. 2019. Microbial biospherics: the experimental study of ecosystem function and evolution. *Proceedings of the National Academy of Sciences of the USA* 116:11093–11098.

Rocha, O. J., and E. Braker. 2021. The Organization for Tropical Studies: history, accomplishments, future directions in education and research, with an emphasis in the contributions to the study of plant reproductive ecology and genetics in tropical ecosystems. *Biological Conservation* 253:108890.

Rushworth, C. A., R. S. Baucom, B. K. Blackman, M. Neiman, M. E. Orive, A. Sethuraman, J. Ware, et al. 2021. Who are we now? a demographic assessment of three evolution societies. *Evolution* 75:208–218.

Sanchez, A., S. Estrela, and M. Rebolledo-Gomez. 2021. Multi-replicated enrichment communities as a model system in microbial ecology. *EcoEvoRxiv*, <https://ecoevrxiv.org/pzh82>.

Schell, C. J., C. Guy, D. S. Shelton, S. C. Campbell-Staton, B. A. Sealey, D. N. Lee, and N. C. Harris. 2020. Recreating Wakanda by promoting Black excellence in ecology and evolution. *Nature Ecology and Evolution* 4:1285–1287.

Sensoy, Ö., and R. DiAngelo. 2017. "We are all for diversity, but . . .": how faculty hiring committees reproduce Whiteness and practical suggestions for how they can change. *Harvard Educational Review* 38:557–580.

Shansky, R. M. 2019. Are hormones a "female problem" for animal research? *Science* 364:825–826.

Stevens, K. R., K. S. Masters, P. I. Imoukhuede, K. A. Haynes, L. A. Setton, E. Cosgriff-Hernandez, M. A. Lediju Bell, et al. 2021. Fund Black scientists. *Cell* 184:561–565. <https://doi.org/10.1016/j.cell.2021.01.011>.

Stewart, A. J., and V. Valian. 2018. An inclusive academy. MIT Press, Cambridge, MA.

Stocks, G., L. Seales, F. Paniagua, E. Maehr, and E. M. Bruna. 2008. The geographical and institutional distribution of ecological research in the tropics: the geography of tropical ecology. *Biotropica* 40:397–404.

Stone, D. E. 1988. The Organization for Tropical Studies (OTS): a success story in graduate training and research. Pages 143–187 in F. Aldemeda and C. M. Pringle, eds. Tropical rainforest diversity and conservation. California Academy of Sciences and Pacific Division, AAAS, San Francisco.

Strong, D. R. 1977. Insect species richness: hispine beetles of *Heliconia latispatha*. *Ecology* 58:573–582.

Thompson, J. E., and T. R. Birkhead. 2020. Avian egg collections: museum collection bias driven by shape and size. *Journal of Avian Biology* 51:e02507.

Tilman, D., P. B. Reich, J. Knops, D. Wedin, T. Mielke, and C. Lehman. 2001. Diversity and productivity in a long-term grassland experiment. *Science* 294:843–845.

Travis, J. 2006. Is it what we know or who we know? choice of organism and robustness of inference in ecology and evolutionary biology. *American Naturalist* 167:303–314.

Tseng, M., R. W. El-Sabaawi, M. B. Kantar, J. H. Pantel, D. S. Srivastava, and J. L. Ware. 2020. Strategies and support for Black, Indigenous, and people of colour in ecology and evolutionary biology. *Nature Ecology and Evolution* 4:1288–1290.

Tydecks, L., V. Bremerich, I. Jentschke, G. E. Likens, and K. Tockner. 2016. Biological field stations: a global infrastructure for research, education, and public engagement. *BioScience* 66:164–171.

Valantine, H. A., P. K. Lund, and A. E. Gammie. 2016. From the NIH: a systems approach to increasing the diversity of the biomedical research workforce. *CBE Life Sciences Education* 15:fe4.

Wale, N., and M. A. Duffy. 2021. The use and underuse of model systems in infectious disease ecology and evolutionary biology. *American Naturalist* 198:69–92.

Wall Kimmerer, R. 2013. Braiding sweetgrass. Milkweed, Minneapolis.

Wanelik, K. M., J. S. Griffin, M. L. Head, F. C. Ingleby, and Z. Lewis. 2020. Breaking barriers? ethnicity and socioeconomic background impact on early career progression in the fields of ecology and evolution. *Ecology and Evolution* 10:6870–6880.

Wehi, P. M., H. Whaanga, and S. A. Trewick. 2012. Artefacts, biology and bias in museum collection research. *Molecular Ecology* 21:3103–3109.

Yang, L. H. 2004. Periodical cicadas as resource pulses in North American forests. *Science* 306:1565–1567.

Zea, M. C., and L. Bowleg. 2016. The final frontier-transitions and sustainability: from mentored to independent research. *AIDS and Behavior* 20(suppl. 2):311–317.

Altermatt, F., E. A. Fronhofer, A. Garnier, A. Giometto, F. Hammes, J. Klecka, D. Legrand, et al. 2015. Big answers from small worlds: a user's guide for protist microcosms as a model system in ecology and evolution. *Methods in Ecology and Evolution* 6:218–231.

Anderson-Teixeira, K. J., S. J. Davies, A. C. Bennett, E. B. Gonzalez-Akre, H. C. Muller-Landau, S. J. Wright, K. Abu Salim, et al. 2015. CTFS-ForestGEO: a worldwide network monitoring forests in an era of global change. *Global Change Biology* 21:528–549.

Bedford, N. L., and H. E. Hoekstra. 2015. *Peromyscus* mice as a model for studying natural variation. *eLife* 4:e06813.

Blount, Z. D. 2015. The unexhausted potential of *E. coli*. *eLife* 4:e06956.

Denell, R. 2008. Establishment of *Tribolium* as a genetic model system and its early contributions to evo-devo. *Genetics* 180:1779–1786.

Duan, S.-F., P.-J. Han, Q.-M. Wang, W.-Q. Liu, J.-Y. Shi, K. Li, X.-L. Zhang, and F.-Y. Bai. 2018. The origin and adaptive evolution of domesticated populations of yeast from Far East Asia. *Nature Communications* 9:2690.

Ebert, D. 2011. A genome for the environment. *Science* 331:539–540.

Frézal, L., and M.-A. Félix. 2015. *C. elegans* outside the petri dish. *eLife* 4:e05849.

Grant, P. R., and B. R. Grant. 2002. Unpredictable evolution in a 30-year study of Darwin's finches. *Science* 296:707–711.

Grant, R. B. 2003. Evolution in Darwin's finches: a review of a study on Isla Daphne Major in the Galápagos Archipelago. *Zoology* 106:255–259.

Griffiths, A. G., R. Moraga, M. Tausen, V. Gupta, T. P. Bilton, M. A. Campbell, R. Ashby, et al. 2019. Breaking free: the genomics of allopolyploidy-facilitated niche expansion in white clover. *Plant Cell* 31:1466–1487.

Jones, F. C., M. G. Grabherr, Y. F. Chan, P. Russell, E. Mauceli, J. Johnson, R. Swofford, et al. 2012. The genomic basis of adaptive evolution in threespine sticklebacks. *Nature* 484:55–61.

Krämer, U. 2015. Planting molecular functions in an ecological context with *Arabidopsis thaliana*. *eLife* 4:e06100.

Kress, W. J., D. L. Erickson, F. A. Jones, N. G. Swenson, R. Perez, O. Sanjur, and E. Birmingham. 2009. Plant DNA barcodes and a community phylogeny of a tropical forest dynamics plot in Panama. *Proceedings of the National Academy of Sciences of the USA* 106:18621–18626.

Lampert, W. 2011. *Daphnia*: development of a model organism in ecology and evolution. International Ecology Institute, Oldendorf.

Lee, B.-Y., B.-S. Choi, M.-S. Kim, J. C. Park, C.-B. Jeong, J. Han, and J.-S. Lee. 2019. The genome of the freshwater water flea *Daphnia magna*: a potential use for freshwater molecular ecotoxicology. *Aquatic Toxicology* 210:69–84.

Lowry, D. B., J. M. Sobel, A. L. Angert, T. Ashman, R. L. Baker, B. K. Blackman, Y. Brandvain, et al. 2019. The case for the continued use of the genus name *Mimulus* for all monkeyflowers. *Taxon* 68:617–623.

Magurran, A. E. 2005. Evolutionary ecology: the Trinidadian guppy. Oxford University Press, Oxford.

Markow, T. A. 2015. The secret lives of *Drosophila* flies. *eLife* 4:e06793.

McKenna, D. D., and B. D. Farrell. 2006. Tropical forests are both evolutionary cradles and museums of leaf beetle diversity. *Proceedings of the National Academy of Sciences of the USA* 103:10947–10951.

McKinnon, J. S., and H. D. Rundle. 2002. Speciation in nature: the threespine stickleback model systems. *Trends in Ecology and Evolution* 17:480–488.

Muschiol, D., F. Schroeder, and W. Traunspurger. 2009. Life cycle and population growth rate of *Caenorhabditis elegans* studied by a new method. *BMC Ecology* 9:14.

Neff, E. P. 2020. Where the wild zebrafish are. *Lab Animal* 49:305–309.

Parichy, D. M. 2015. Advancing biology through a deeper understanding of zebrafish ecology and evolution. *eLife* 4:e05635.

Park, T. 1948. Experimental studies of interspecies competition. I. Competition between populations of the flour beetles, *Tribolium confusum* Duvall and *Tribolium castaneum* Herbst. *Ecological Monographs* 18:267–307.

Phifer-Rixey, M., and M. W. Nachman. 2015. Insights into mammalian biology from the wild house mouse *Mus musculus*. *eLife* 4:e05959.

Replansky, T., V. Koufopanou, D. Greig, and G. Bell. 2008. *Saccharomyces* sensu stricto as a model system for evolution and ecology. *Trends in Ecology and Evolution* 23:494–501.

Rojas, B., E. Burdfield-Steel, C. De Pasqual, S. Gordon, L. Hernández, J. Mappes, O. Nokelainen, et al. 2018. Multimodal aposematic signals and their emerging role in mate attraction. *Frontiers in Ecology and Evolution* 6:93.

Tribolium Genome Sequencing Consortium. 2008. The genome of the model beetle and pest *Tribolium castaneum*. *Nature* 452:949–955.

US Fish and Wildlife Service. 2018. Zebra danio (*Danio rerio*): ecological risk screening summary. <https://www.fws.gov/fishes/ans/erss/uncertainrisk/ERSS-Danio-rerio-FINAL-July2018.pdf>.

Wilf, P., C. C. Labandeira, W. J. Kress, C. L. Staines, D. M. Windsor, A. L. Allen, and K. R. Johnson. 2000. Timing the radiations of leaf beetles: hispines on gingers from latest cretaceous to recent. *Science* 289:291–294.

Wu, C. A., D. B. Lowry, A. M. Cooley, K. M. Wright, Y. W. Lee, and J. H. Willis. 2008. *Mimulus* is an emerging model system for the integration of ecological and genomic studies. *Heredity* 100:220–230.

Yen, E. C., S. A. McCarthy, J. A. Galarza, T. N. Generalovic, S. Pelan, P. Nguyen, J. I. Meier, et al. 2020. A haplotype-resolved, de novo genome assembly for the wood tiger moth (*Arctia plantaginis*) through trio binning. *GigaScience* 9:giaa088.

Zakhartsev, M., and M. Reuss. 2018. Cell size and morphological properties of yeast *Saccharomyces cerevisiae* in relation to growth temperature. *FEMS Yeast Research* 18:foy052. <https://doi.org/10.1093/femsyr/foy052>.

Zalucki, M. P., and A. R. Clarke. 2004. Monarchs across the Pacific: the Columbus hypothesis revisited. *Biological Journal of the Linnean Society* 82:111–121.

Zhan, S., W. Zhang, K. Niitsepöld, J. Hsu, J. F. Haeger, M. P. Zalucki, S. Altizer, et al. 2014. The genetics of monarch butterfly migration and warning colouration. *Nature* 514:317–321.

Editor: Daniel I. Bolnick



“*Trochilus colubris* taking food, drawn from memory.” From “Ornithophilous Pollination” by Joseph L. Hancock (*The American Naturalist*, 1894, 28:679–683).