Niche perspectives on plant-pollinator interactions

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Ryan D. Phillips^{1,2,3}*, Rod Peakall³, Timotheüs van der Niet⁴, and Steven D. Johnson⁴

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- ¹Department of Ecology, Environment and Evolution, La Trobe University, Vic.
- 6 3086, Australia.
- ²Kings Park Science, Department of Biodiversity, Conservation and Attractions, WA
- 8 6005,
- 9 Australia.
- ³Ecology and Evolution, Research School of Biology, The Australian National
- 11 University, Canberra, ACT 0200, Australia.
- ⁴Centre for Functional Biodiversity, School of Life Sciences, University of KwaZulu-
- Natal, P Bag X01, Scottsville 3209, South Africa.
- *Correspondence: R.Phillips@latrobe.edu.au

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Abstract

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- 20 While ecological niches are critical to species co-existence and diversification, the
- 21 niche concept has been under-utilized in studying the roles of pollinators in plant
- evolution and reproduction. Pollination niches can be objectively characterized using
- pollinator traits, abundance and distributions, as well as network topology. We review
- evidence that floral traits represent adaptations to pollination niches, with trade-offs in
- trait deployment reinforcing niche specialisation. In turn, specialised pollination
- 26 niches potentially increase speciation rates, foster species co-existence and constrain
- species range limits. By linking studies of adaptation with those on speciation and co-
- existence, the pollination niche provides an organising principle for research on plant
- reproduction and conceptually unites these studies with fields of biology where the
- niche perspective is already firmly established.

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The importance of the pollination niche

Understanding the diversification of flowering plants remains one of the great 33 challenges in biology [1]. Fundamental to understanding the processes underpinning 34 diversification is the concept of the **niche** (see Glossary), the combination of resources and environmental conditions that enable a population to establish positive 36 growth rates if individuals possess appropriate functional traits [2]. Niches are an important driver of trait evolution, potentially leading to the origin of new species, but are also critical for determining species' geographic range. Partitioning of niches also reduces competition and is thus an important contributor to species co-existence [2, 3]. Resolving the functional traits associated with niche occupancy gives the niche concept tremendous potential to integrate studies of trait adaptation and community ecology [4]. While discussion of niches in plants has tended to focus on abiotic factors, as most plant species are reliant on pollination for reproduction and therefore population persistence, pollination is also a critical component of a plant's niche [5]. Using the concept of the **pollination niche**, we develop a framework that unites studies of floral adaptation, species co-existence, plant distribution and speciation,

with the overall goal of understanding the origins and maintenance of diversity of flowering plants. Having defined the pollination niche, we discuss approaches that can be used for quantifying pollination niches, including potential opportunities for using plant-pollinator network methodologies, before addressing the following questions: (i) can the pollination niche act as a basis to understand functional traits? (ii) do **trade-offs** explain occupancy of specialised pollination niches? (iii) what is the role of the pollination niche in shaping community structure? (iv) to what extent does the pollination niche determine geographic ranges? (v) do certain key floral innovations increase speciation rates by enabling occupation of diverse pollination niches?

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What is the pollination niche?

We consider the **fundamental pollination niche** to be the vectors (biotic or abiotic) that can potentially enable pollination for a given plant species, noting that the fundamental niches at any given site may be expected to reflect the local temporal and spatial variability of the pollinator assemblage. The **realised pollination niche** is shaped by community level interactions affecting the effectiveness of pollen vectors due to both the competition for pollinators among plants and the foraging preferences

of pollinators. Not all pollination niches in a community may be occupied at a given point in time. For example, many species of hymenopteran use pheromone systems to attract mates, yet only a tiny fraction of these insects are exploited for pollination by plants that mimic insect sex pheromones [6]. Similarly, many generalist birds will consume nectar if it is available, yet these birds are exploited for pollination only in certain geographical regions [7].

The presence of a group of pollinators in a plant community and their relative abundance clearly represents the most critical aspect of the pollination niche. However, other axes of the pollination niche include seasonal opportunities for pollination [8], variation in the lengths of the mouthparts with which pollinators interact with flowers [9, 10], sites of pollen deposition on the pollinator [11-13], diel opportunities for pollen release or pollinator visitation [14], and the type of floral reward sought by animals [15, 16]. Each of these different niche axes has the potential to drive floral trait diversification [17] and reduce interspecific pollen transfer [18], both factors potentially reducing interspecific competition and maintaining reproductive isolation.

Quantifying pollination niches

For pollination niches to be a useful tool for understanding the origins of plant diversity, an objective approach for recognising and defining them is needed. In their simplest form, niches can be described by niche position and niche breadth along an axis. For example, in terms of a plant's edaphic preferences, there will be an optimal soil pH for performance (niche position), but also a range of pH where population growth rates are positive (niche breadth). When multiple variables are considered, the niche becomes the area in multi-dimensional space where growth rates are positive [19].

With regards to pollination, the niche would be the set of animal species or functional group(s) and their associated traits that lead to positive population growth rates. For example, for a hawkmoth-pollinated plant species, the tongue length of hawkmoth pollinators can be used to identify niche position (median tongue length) and niche breadth (range of tongue lengths for effective pollination) [9]. This approach can be extended to the full plant community across multiple pollinators and their traits to

quantify areas of niche space that plants could potentially occupy (**Box 1**). However, it is important to note that for those plant species reliant on biotic vectors, not all visitors are part of the realised niche. For example, in a plant adapted to pollination by sunbirds, the niche position critical for population persistence would include sunbirds with particular bill lengths, but not incidental insect visitors that are ineffective pollinators [20, 21].

There is a vast taxonomic and functional diversity of animals involved in pollination, and for many of these we lack details of their behavioural responses to flowers and their effect on plant fitness. As such, in pollination biology, the concept of niche position is often expressed qualitatively in terms of pollinator functional groups - a group of pollinators that are likely to exert similar selection pressures [16, 22] (most often taxonomically related e.g. hawkmoths, sunbirds). The use of pollinator functional groups is an effective tool in ecological and evolutionary studies when large differences in the morphology and/or behaviour of the pollinators make the pollinator niches relatively discrete. However, the inherently nested nature of pollinator functional groups can make comparison between species or communities challenging [5]. For example, sunbirds are nested within the broader category of bird pollinators, but among the sunbirds there is also variation in beak length among species [23]. Thus, a reliance on conventionally recognised functional groups may completely overlook novel or subtle groupings leading to an underestimate of niche specialisation. Therefore, visitation data for entire plant-pollinator networks could provide a technique for more objective recognition of niches and the associated level of specialisation.

Are network approaches a suitable first step for identifying pollination niches?

Recognising pollination niches using plant-pollinator networks

Plant-pollinator networks describe the patterns of pollinator visits to plants for a community of plants and pollinators (e.g. [24-27]). By characterising plant-pollinator interactions at the community level, networks reveal not only interacting partners, but also which potential interactions are absent among the species that are present in the community. This has the potential to help researchers characterize the realised pollination niche of individual species relative to the total pool of pollination niches available. Further, analysing patterns of visitation for the entire community avoids

potential biases about which species are candidates for the use of particular niches, and may even help to identify niches not previously suspected to exist.

The detection of **modularity**, where groups of species interact more frequently with each other than with the remainder of the network, might offer an important first step toward objectively identifying pollinator functional groups (niche position) and relatively discrete pollination niches [27-29] **Figure 1**). While it can be tested if entire networks exhibit significant modularity (see [28, 30] for methodology), recognition of particular pollination niches requires identification of the specific modules to which the plant and pollinator species have been assigned [27, 29]. Importantly, since networks of plant-pollinator interactions are often nested, analysis of modularity can be undertaken in a hierarchical fashion to test for modules within modules (e.g. long-billed birds may be nested within a bird pollination module). However, given that modularity is a consistent structural trait of plant-pollinator communities, and that it is more likely to be detected in larger networks [28, 30], additional evidence may be needed to test whether the modules identified are both biologically meaningful and correspond to niches.

As a first step for testing the functional significance of modules, the plant members of a given module must be shown to be interacting with any shared pollinators via the same general pollination mechanism (e.g. exploitation of nectar foraging versus sexually deceived insects). This is necessary to avoid inadvertently lumping plants that actually occupy different niches. Once the basis of the interaction is known to be consistent within a module, one can then test the expectation that pollinators from within modules will be more effective than pollinators from other modules. If this prediction is supported, and the module is indicative of a particular pollination niche, one can test for an association between the module and specific floral traits. Finally, testing functional significance of modules might involve experimentally establishing whether the pollinators assigned to a module show similar responses to the flowers, and if floral traits associated with the module influence pollinator attraction. As such, the identification of network modules could represent a critical step towards testing hypotheses to explain floral adaptation as well as community processes [24, 25, 27].

Plant-pollinator networks and specialisation

Since the study of interaction networks has become widely employed in pollination biology, understanding specialisation has been a common topic of interest for researchers taking this approach (see references in [31]). However, there are several practical and statistical challenges to effectively implementing network studies, with important consequences for quantifying the specialisation of pollination niches (Box 2). For example, absolute ecological specialisation cannot simply be measured by metrics such as k (i.e. number of links - the number of other species that a given species interacts with), as they partly reflect sampling effort [32]. Recognition of this bias lead to the widespread usage of complementary specialisation d' [33], which describes the deviation from a neutral pattern of associations [34]. However, rather than down-weighting the specialisation of a species because it uses an abundant pollinator, as in d, we believe a more rigorous approach is to test whether the species shows ecological specialisation and to identify the mechanism underpinning this specialisation. Ecological specialisation is best quantified by identifying effective pollinator species using methods such as single-visit pollen deposition and selective exclusion experiments [35], and then using rarefaction approaches [36] to estimate the level of sampling required to gain an accurate estimate of the number of interaction partners [37]. Further, the proportion of pollination events by the most frequent of the effective pollinator species, as an estimate of the reliance on a single species, is potentially a useful metric for studies in both ecology and evolution. Once absolute ecological specialisation is estimated, further data can be gathered to address whether the plant attracts particular pollinators via specialised signals or rewards that are adaptations to particular pollinators (i.e. is phenotypically specialised) or if specialisation is merely imposed by an absence of other potentially suitable pollinators [22].

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Functional floral traits

In plant community ecology, functional traits, such as seed size, canopy height and leaf mass per area, have been considered to represent adaptations to particular environmental conditions or life history strategies, [4, 38]. Functional traits have the ability to determine the structure of plant communities by imposing a mechanism that filters out species that are not suitably equipped to occupy any of the available niches, or are outcompeted by species already occupying certain niches. Functional traits therefore ultimately affect which niches species can exploit and whether they can co-

exist [39]. For the best known pollination groups, it is clear that traits such as floral colour, odour and morphology can be considered functional traits where certain character states provide a fitness advantage for the occupation of particular pollinator niches (Table 1). Manipulation of floral traits has demonstrated that divergence in floral traits is associated with differences in pollinator attraction and plant fitness [17, 40-43]. Further, experimental studies have confirmed that heritable changes in floral traits can occur in just a few generations when plants are exposed to different pollinator environments [44, 45]. At the macroecological level, studies analysing pollinator data and floral traits simultaneously in a phylogenetic context have provided compelling evidence for associations between particular pollinators and sets of floral traits [10, 46, 47]. For example, there has been convergent evolution of red floral colouration and, in some cases large nectar quantities, exserted anthers and changes in corolla shape and spur length, associated with shifts from bee to hummingbird pollination in several plant genera in the North American flora such as Penstemon, Mimulus and Aquilegia [10, 46, 48, 49]. Floral functional traits may offer the same, or even better, predictive power for

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assessing niche occupancy than do the vegetative traits that are traditionally used in plant community ecology [4, 38]. This is evidenced by the extensive literature on pollination syndromes that, despite being developed largely independently of niche theory, has demonstrated the association of certain floral colours, shapes and rewards with particular functional groups of pollinators [5, 47, 50-52]. For example, a recent analysis of the South African flora, which focused on species with putatively specialised pollination systems, showed that there is a close correspondence (c. 80%) matching) between Stefan Vogel's [51] assignments of plant species to floral syndromes in the 1950s and the subsequently discovered pollinators of these species [53]. However, support for pollination syndromes in some other floras, particularly those in northern temperate regions, is more equivocal [54], suggesting that the concept has less utility when pollination systems are relatively generalized. Pollination syndromes may also be obscured when floral traits more strongly reflect the bauplan of the plant lineage than adaptations to pollinators. In addition, most floral syndrome classifications are based on qualitative trait categories, including subjective human assessments of colour and scent. As such, there is a need for floral

functional traits to be standardized in the same quantitative manner as plant vegetative traits [38] and validated across geographical regions and plant groups.

Do trade-offs explain niche specialisation?

If visitation from multiple pollinators (niche generalization) has the potential to provide greater and more stable pollination service, it raises the question of how and why niche specialisation arises? One potential explanation is that adaptation to the most effective pollinator(s) involves trade-offs, where the functional traits that increase visitation or effectiveness of one pollinator species reduce attraction of other potential pollinator species or their pollination effectiveness [55]. Trade-offs in plants are well known for traits related to survival, such as the trade-off between growth rate and nutrient retention that leads to differences in growth rate between species occurring in high and low nutrient conditions [56]. In pollination systems, such trade-offs could act at multiple stages through the pollination process including long-distance attraction to floral signals, behaviour on the flower, and morphological fit. One obvious implication of strong-trade-offs is that hybrids between species with different specialised pollination systems will often have reduced pollination success, thus resulting in at least a partial barrier to gene flow between species.

The strongest evidence for the presence of trade-offs underpinning specialised pollination niches comes from systems where divergent floral traits associated with the attraction of two pollinator species are manipulated to test whether a change to intermediate floral traits is associated with reduced pollinator attraction or effectiveness. For example, trade-offs have been demonstrated for floral signals involved in pollinator attraction [42] and in floral traits affecting the morphological fit of the pollinator [57] (**Figure 2**). However, **bimodal pollination** has been reported for some plant species (e.g. involving mammals and hummingbirds with parallel adaptations to both groups [58]) and many plants have relatively generalist pollination systems, suggesting that trade-offs are not always strong enough to enforce specialisation. Nonetheless, the few experimental tests of trade-offs have generally provided strong evidence in support of their existence [41, 42, 57, 59], but see [60, 61]. Should future experiments confirm that trade-offs are a key mechanism underpinning specialisation, this would also support the hypothesis that trait trade-offs

contribute to the process of speciation and to the persistence of diverse plant communities.

Pollination niches and species co-existence

The occupation of different niches as a mechanism that alleviates interspecific competition is often critical for species co-existence [3]. For example, in plants, different drought and water-logging tolerance can lead to the exploitation of different microsites, thereby fostering co-occurrence [19, 62]. Given the large diversity of co-flowering plant species in many plant communities, it is expected that at least some level of partitioning of the available pool of potential pollinators between plants occurs through the use of different pollinator species, thereby reducing interspecific pollen transfer [18] and competition for pollinators [63]. Indeed, in a fascinating recent example, it was shown that two colour morphs of a bromeliad persist through different foraging preferences between males and females of a hummingbird pollinator species, thus preventing competitive exclusion of one of the morphs [64].

Pollination niches are unusual in the niche context for not only providing scope for interspecific competition, but also for ecological facilitation. Rewarding plants often exhibit Allee effects where small populations produce a lower total reward to pollinators and therefore achieve lower per plant visitation [65-67]. A corollary of this is that a plant can benefit from the presence of other co-occurring rewarding "magnet" plant species that attract a greater number of pollinators into the plant community because of an effective increase in the total number of rewarding plants. While perhaps best known in deceptive orchids, where non-rewarding plants receive a benefit from co-occurring with rewarding plants [68, 69], facilitation by attracting additional pollinators has also been demonstrated among different rewarding plant species [70, 71]. However, experiments with the self-incompatible herb *Raphanus raphanistrum* demonstrated the context dependence of such effects [72]. Using experimental plots with plants that shared pollinator species, it was shown that facilitative effects transitioned to competitive effects as the abundance of the focal species was reduced relative to the co-occurring plant species.

While using different pollinators clearly reduces the potential for competition between plants, the question remains whether partitioning of the pollination niche is a major

contributor to co-existence of closely related or ecological similar plants. Pauw [63] hypothesised that when reproduction is pollen limited and the abundance of a plant species is limited by intraspecific competition for pollinators, a new colonist (or mutation) in the community is more likely to establish if it occupies a different pollination niche. Using a model of plant and pollinator dynamics, Benadi [73] showed that the occupation of different pollination niches is most likely to contribute to species co-existence when the costs of searching for flowers are low, pollinator populations are strongly limited by resources other than floral rewards, and plant-pollinator interactions are specialised. There is some empirical evidence that the rate of per flower pollinator visits declines at both low and particularly high levels of plant abundance [74, 75], suggesting the potential for competition between plant species that share pollinators [76].

Several community studies have tested if some plant communities are structured such that there is much lower pollination niche overlap than expected by chance. For example, there is evidence of interspecific competition being avoided by differential pollen placement [11, 13], the use of different pollinator species [12, 77], and differences in pollen release [14] and flowering time [8]. However, there are also several studies that have not found any strong signature of competition for pollinators in terms of community structure [78]. Interestingly, this includes research on the coexistence of species in the hyper-diverse assemblages of *Erica* in the Cape Floristic Region of South Africa where evidence of strong niche partitioning was anticipated [79, 80]. An unresolved issue in systems where competition does appear to be important, is whether competition mainly results in *in situ* **character displacement** or filtering (ecological sorting) of plants that can enter the community.

The pollination niche and species range limits

While the spatial distribution of plants is strongly correlated with edaphic and climatic niche components [81, 82], pollinators also have the ability to limit the spatial distribution of plants. Indeed, at broader spatial scales there is evidence that the pollination niche can limit the geographic range of plant species [83]. Using niche modelling, Duffy & Johnson [84] showed that for 21 of 32 South African plant species with relatively specialised pollination systems, pollinator distribution was the best predictor of their geographic range. Though pollinator constraints on plant

geographic ranges would intuitively be more likely to apply in plants with specialised niches, one of the most compelling cases of pollinators contributing to natural range limits comes from a plant species pollinated by several generalist bee species. In the Californian endemic *Clarkia xantiana* the abundance of generalist bees is lower and more variable at the more arid range margin leading to overall lower pollinator availability [85]. Additional evidence for pollinator limitation on geographic ranges comes from anthropogenic dispersal of plants, where of the 1000s of plant introductions, there are few cases of plants with highly ecologically specialised pollination systems becoming invasive [86, 87]. For example, of the 60 species of *Ficus* introduced to Florida, beyond the natural geographic range of their specific pollinator species, the only three to become invasive did so after the introduction of their pollinator [86]. Similar processes appear to operate at the landscape scale, where scarcity of pollinator species can lead to low levels of reproduction, potentially leading to reduced occupancy of otherwise suitable patches of habitat [88, 89].

The pollination niche and evolutionary diversification

Pollination has long been hypothesised to be a key contributor to the diversification of flowering plants [90-92]. Indeed, there is abundant evidence that pollinators drive phenotypic diversification among populations [92]. Due to the link between pollinator adaptation and the evolution of reproductive isolation, these microevolutionary processes are expected to result in a macroevolutionary pattern of adaptive radiations in pollination systems [5, 47, 93]. The niche perspective is central for understanding the role of pollination in plant diversification at two levels. Firstly, the evolution of a particular pollination strategy or even a single floral trait may greatly increase the number of niches available for exploitation, which may, in turn, promote diversity of the lineage. As such, key innovations such as zygomorphy [94] or nectar spurs [95] have been linked to increased diversification rate. The process underlying this increased rate is thought to be the exploitation of an increased number of relatively discrete pollination niches, but this has not been tested in detail. Key innovations can also include chemical traits; for example, the evolution of pollination by sexual deception can lead to a large number of new pollination niches, each involving exploitation of the males of particular insect species [6, 96], potentially leading to an overall increase in plant diversification rate [97]. Secondly, some pollination niches may be indirectly associated with elevated levels of plant speciation. For example, it

has been predicted that in some plant groups diversification rates will be higher in bee-pollinated than in bird-pollinated lineages due to birds reducing genetic differentiation through long-distance pollen dispersal [49]. In some lineages and some geographic regions there is compelling evidence that shifts in the pollination niche are often associated with speciation. Indeed, in an analysis of plant lineages where phylogenetic and pollination strategy data are available, Van der Niet & Johnson [93] showed that approximately 25% of speciation events were associated with a shift in pollinator type. However, it remains poorly understood how frequently shifts in pollinator assemblages are associated with speciation in generalist lineages, and if any such differences arise from evolution of floral traits or as a simple by-product of related plant species occupying habitats or geographic regions with different pollinator assemblages [98, 99].

Concluding remarks and future perspectives

Studies of plant-pollinator interactions are typically conducted according to two very different traditions: classical adaptationist studies [17] and community-level network studies [31]. There has been little cross-talk between these two approaches, with the adaptationist approach emphasizing trait functions and factors that impact on plant fitness, and the network approach emphasizing metrics that describe the ecological properties of whole communities. The niche perspective has potential to integrate these two approaches to help explain the community context of specialisation in pollination systems and its implications for trait evolution, lineage diversification, and species persistence. Likewise, the evolution of floral traits in generalist lineages could potentially be explained through using community data for the complex task of defining generalist niches, and by experiments that test for variation between pollinator species in both their response to floral traits and their consequences for plant fitness [61].

While objective recognition of niches is becoming a cornerstone of studies using the pollination niche concept, there is a need to expand beyond using only network summary analyses of patterns of plant-pollinator interactions at a particular point in space and time. For example, experimental manipulation of the flowering or pollinating community (e.g. [100-102]) is a potentially powerful, but currently underutilised, approach to improve understanding the fundamental niche.

Incorporating morphology and pollinator behaviour into studies of interaction 405 networks enables a direct test of whether floral traits are associated with network 406 modules (e.g. [24]), thereby identifying traits that can be tested experimentally for 407 their role in pollinator attraction (see Outstanding Questions). The identification of 408 functional traits is critical as this forms the basis of understanding specialisation and 409 how shifts in niches may occur - one of the most fundamental unanswered questions 410 in pollination biology. Indeed, the evidence for numerous shifts in pollination strategy 411 in many plant lineages [93] highlights the importance of understanding how transitions arise between pollination niches. A key aim for future studies will be to 413 identify the pollination niche axes that are most often associated with plant speciation 414 and to determine whether these are the same or different from the axes that are most 415 effective at fostering ecological co-existence. We expect that the incorporation of 416 niche theory into pollination biology will enable much better integration of this 417 discipline into the general fields of ecology and evolutionary biology. 418

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684	Glossary:
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- Bimodal pollination: where for a given plant species pollination is achieved by two
- different functional groups of animals normally associated with different sets of
- 687 functional traits.
- 688 **Character displacement:** evolution through natural selection for traits that reduce
- competition or reproductive interference.
- **Ecological specialisation:** in the context of pollination, the use of one or few
- 691 pollinator species.
- Functional trait: a trait that strongly affects organismal performance. As such, they
- are expected to have a large affect on fitness in any given niche.
- **Fundamental niche:** In the context of biotic pollination, the pollen vectors capable of
- supporting positive population growth rates in the absence of competition.
- 696 **Modularity:** A consistent pattern in interaction networks where groups of species
- tend to interact with each other more regularly than with the members of other groups.
- Niche: the resources and environmental conditions that enable a population of a
- species to maintain a net reproductive rate of ≥ 1 . Each variable contributing towards
- the overall niche is referred to as a niche axis. The niche of a species is commonly
- articulated as the n-dimensional hyper-volume generated from the suitable conditions
- n across n niche axes.
- Plant-pollinator network: the set of pollination interactions in a community of
- 704 plants and pollinators.
- Pollinator functional group: a group of pollinator species that are likely to exert
- similar selection pressures. These pollinator species are often phylogenetically
- 707 related.
- Pollination niche: the abiotic or biotic vectors and their associated traits providing
- effective pollination at the time and place that a plant species flowers.
- Pollination syndrome: a suite of floral traits shared across plant species that
- primarily use the same pollinator functional group.
- Realised niche: In the context of biotic pollination, the pollination service from the
- pollen vectors that can support positive population growth rates, as shaped by
- 714 competition and facilitation.
- **Trade-offs:** when adaptation to a particular niche lowers performance in another
- 716 niche.

Box 1. Using pollinator traits to recognise pollination niches 719 720 Overview 721 Plotting the frequency distribution of variation in hawkmoth tongue length across a 722 pollinator community can aid identification of the potential pollination niches 723 available to the plant species pollinated by nectar foraging hawkmoths in a given 724 community. Subsequently, an independent assessment of plant functional traits 725 associated with hawkmoth pollination, such as flower tube length, can reveal if plants 726 show adaptation to a particular pollination niche [9]. 727 728 Pollination niche position, breadth and availability 729 In this example, pollination niche position is the median tongue length for a given 730 functional pollinator group (Figure I, panel C and D). Pollination niche breadth is the 731 732 range of effective tongue lengths (panel C and D). Pollination niche availability at a given site is indicated by the abundance of pollinator individuals (panel D). In this 733 community, there are two pollination niches, corresponding to short-tongued and 734 long-tongued hawkmoths. Furthermore, the moth-pollinated plant species in the 735 community show strong evidence for floral tube length adaptation to one or other of 736 the two pollination niches (panel E). 737 738 Advantages of using pollinator traits to recognise pollination niches 739 This approach allows one to test for floral adaptation to particular niches, and enables 740 development of clear predictions of niche availability to understand niche occupancy. 741 While illustrated here with a univariate pollinator trait, this approach can be extended 742 to include multiple niche axes in the same way as multi-dimensional quantifications 743 of abiotic niches. For example, other pollinator traits could include food preference, 744 flight time, body size, and floral odour preference. 745 746 Challenges for testing plant adaptation to the pollination niche 747 A priori evidence is needed to identify niche axes that may represent functional traits. 748 Many plant traits that may be adaptations to their pollination niche are multi-variate 749 (e.g. nectar composition) and may require variable reduction methods (e.g. Principal 750 Co-ordinates Analysis), making clear ecological interpretations more difficult. 751

Detailed sampling of the functional group of pollinators, either to determine

- availability or to measure pollinator traits potentially relevant for pollination, can be
- 754 challenging.

Box 2. Challenges for using networks to understand pollination niches. 756 Plant-pollinator networks may allow objective quantification of the pollination niches 757 in any given community. For a given plant species, niche position would be the 758 module, with the discreteness of the niche measured by Modularity Q [30], and niche 759 breadth would be the number of links, or the number of species with more than N 760 number of visits (to remove incidental visitors). Niche availability would be the 761 abundance of effective pollinator individuals summed across species within a module. 762 763 Visitation vs pollination: Quantifying visitation rather than pollination underestimates 764 specialisation of the pollination systems [103, 104]. This has been partly addressed by 765 quantifying contact with the reproductive structures, but evidence suggests that this 766 still can underestimate specialisation compared with quantifying pollen deposition 767 [105]. 768 Detecting pollinators with low visitation rates: Studies that conduct observations 769 based purely on transects rather than equal observation time across plant species bias 770 771 the data towards lower visitation and increased specialisation for rarer plant species [34]. 772 Bias of metrics towards specialisation: Using simulations of networks generated 773 774 under a quantitative niche model, Frund et al. [32] found that 48 metrics of specialisation and network structure are biased towards overestimating specialisation, 775 and advocated at least 20 observations per plant species to accurately estimate 776 specialisation. 777 d' does not measure ecological specialisation: Ecological specialisation of a 778 pollination system is the number of species that an organism interacts with (or 779 primarily with). However, d', which provides a measure of the complementarity (or 780 exclusiveness) of species interactions [34], is frequently used as a measure of 781 ecological specialisation. Pauw & Stanaway [26] illustrated the shortcoming of this 782 approach in a community where a specialist Gladiolus species visited by one species 783 of Anthophora bee species had a d' value of only 0.3 (scale from 0 to 1) while a 784 generalist Crassula species with 10 pollinator species had a d' = 0.99, because nine of 785 its pollinators were not recorded on other plant species. 786 Geography of specialisation: If pooling data across sites, species absent from a site 787 are interpreted as not interacting with particular species when in fact there was no 788

opportunity to interact (forbidden links), resulting in erroneously high specialisation 789 estimates. 790 Temporal/seasonal variation: Networks can change properties over the course of a 791 year and/or flowering season [106]. 792 Pollinators can be attracted via different mechanisms: Not accounting for the 793 mechanism of attraction underestimates the number of pollination niches as the same 794 pollinator could be attracted to two different species through two different 795 rewards/mechanisms. 796

798 Figures

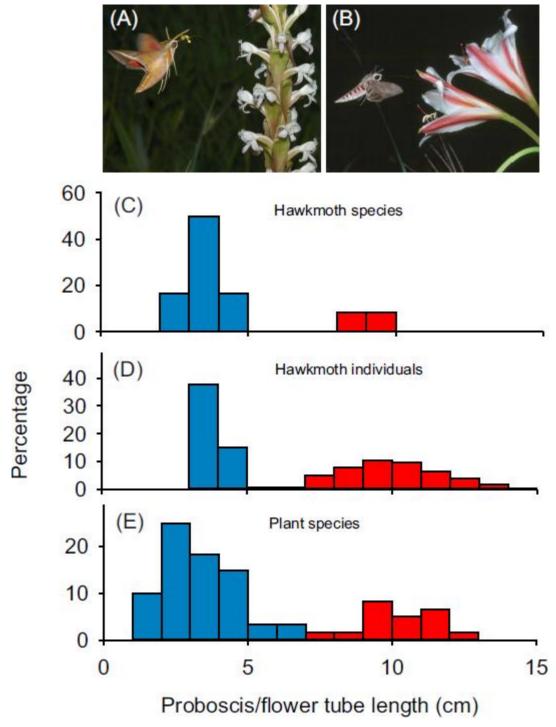


Figure I, Box 1: Subtropical South Africa has numerous species of plant pollinated by hawkmoths, including (A) *Satyrium longicauda* pollinated by the short-tongued hawkmoth *Basiothia schenki* and (B) *Crinum bulbispermum* pollinated by the longtongued hawkmoth *Agrius convolvuli*. Plots illustrate the correspondence between hawkmoth tongue lengths [(C) species means; (D) individual measurements)] and (E) mean floral tube lengths [9]. Blue bars in the histogram illustrate the niche involving

short-tongued hawkmoths, red bars illustrate the niche involving long-tongued hawkmoths. Photos by Steve Johnson.



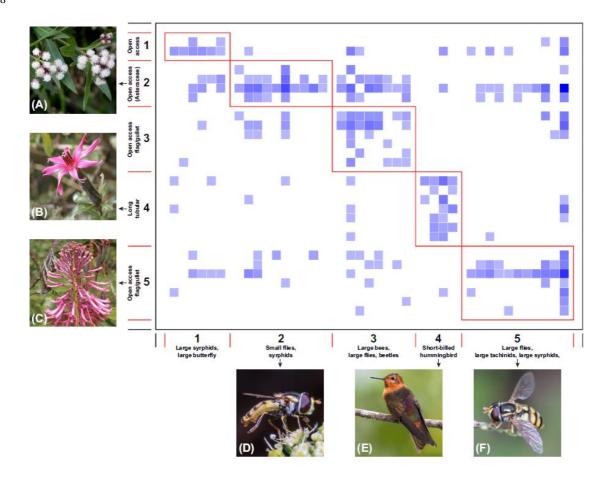


Figure 1: The use of a plant-pollinator network to objectively test for pollinator niches. Potential pollination niches are recognised through the identification of modules, which are then overlayed with data on floral and pollinator traits to test for adaptation by plants to groups of pollinators, and the identification of candidate functional traits. In this example, Watts *et al.* [27] collected data on potential pollinators for several communities in the Peruvian Andes and identified modules using the software *QuantBiMo* [30]. This figure illustrates the objective identification of a module of hummingbirds (e.g. Shining Sunbeam) associated with long tubed flowers (e.g. *Fuchsia apetala*). Other modules mostly correspond to varying groups of insects, particularly flies. This dataset illustrates that some plant species attract pollinators from more than one module, and therefore the potential for individual pollinator species to contribute to more than one pollination niche. For a given plant species, comparing the effectiveness of floral visitors within its module with visitors

from other modules (e.g. for module 4, short-billed hummingbirds vs syrphid flies) could provide a basis for understanding specialisation of the pollination niche. Further work would involve experimental approaches to test if members of these modules respond similarly to floral traits, and if there is evidence for corresponding floral adaptation. Members of modules illustrated are: (A) *Baccaris salcifolia* (Asteraceae); (B) *Barnadesia horrida* (Asteraceae); (C) *Oreocallis grandiflora* (Proteaceae); (D) Syphid species (E) *Aglaeactis cupripennis* (Trochilidae) (F) Syphid species. Note that the syrphid photos are illustrative and do not represent actual members of this community. Photographs: (A) Lynn Watson, (B & C) Stella Watts, (D & F) Thomas Semple, (E) Supreet Sahoo.

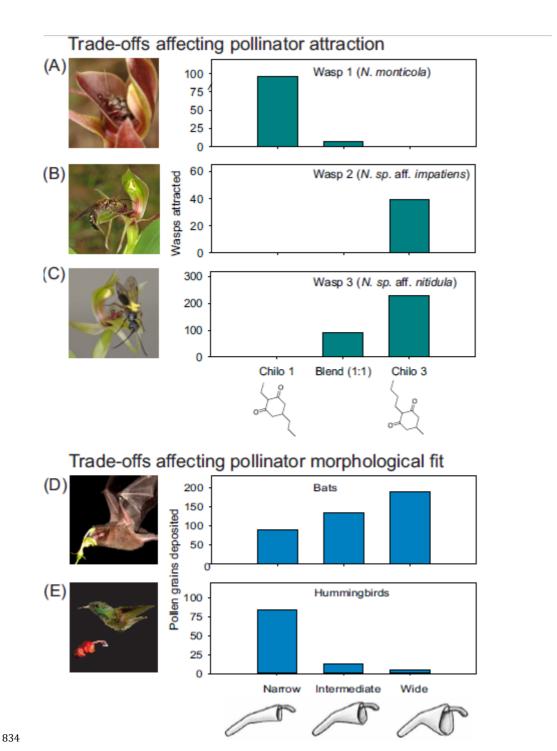
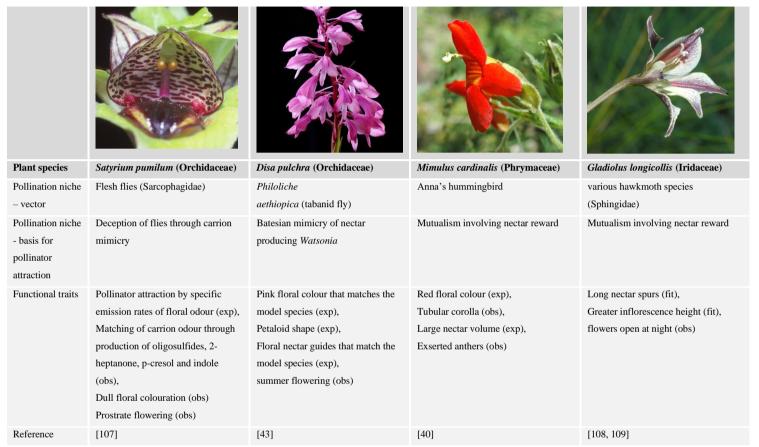


Figure 2: Fitness trade-offs as a mechanism underpinning the specialisation of pollination systems. *Trade-offs in pollinator attraction* (Panel A, B and C): In sexually deceptive orchids, each orchid species is primarily pollinated by a single pollinator species [42, 96], which is attracted through mimicry of the sex pheromone of the female of the pollinator species [6]. Related orchids typically attract related pollinators using different blends of compounds, often with some overlap of the constituent compounds [42]. Here, three species of *Chiloglottis* use one of two different chiloglottones (chiloglottone 1 and 3) to attract sexually deceived

Neozeleboria thynnine wasps. When these compounds are experimentally presented to pollinators either by themselves or in a 1:1 blend, the single compounds attract individuals of different pollinator species, while the blend attracts few or no pollinators (Panel A, B and C) (data from [42]). Trade-offs in morphological fit: Some neotropical plant genera have undergone shifts in pollination strategy between pollination by hummingbirds and bats [47]. Using flowers with artificial corollas in a flight cage experiment, Muchhala [57] demonstrated that narrow corollas guided bills of the hummingbird Adelomyia melanogenys (illustrated here by a similar hummingbird species) more effectively than did wide corollas, leading to greater pollen deposition (Panel E). Alternatively, wide corollas guided snouts of the bat Anoura geoffroyi more effectively than did narrow corollas, leading to more pollen deposition from bats, but less from hummingbirds (Panel D). A model based on this experiment predicted that intermediate corollas are never favoured. Photos by Rod Peakall, Nathan Muchhala and Steven Johnson.

Table 1: Examples of pollination niches and functional traits related to niche occupancy^a



^aFor functional traits; (exp) denotes when the importance of the trait has been tested experimentally for that species; (fit) denotes when variation in plant fitness in response to the trait has been measured in wild populations; (obs) denotes when the importance of the trait has been inferred through observation (e.g. observation of nectar consumption).