Title:

Jack-of-all-trades paradigm meets long-term data: generalist herbivores are more

widespread and locally less abundant

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Abstract

Some of the most common ecological interactions are between plants and herbivorous insects, and these relationships are central to the study of ecological specialization. We address established assumptions about the positive association between local abundance and dietary specialization using a 17-year dataset on the caterpillars of Ecuador. Our long-term data include standardized plot-based samples as well as general, regional collections, allowing for investigations across spatial scales and using different indices of abundance for 1917 morphospecies of Lepidoptera from 33 families. We find that specialists are locally more abundant than generalists, consistent with a key component of the "jack of all trades, master of none" hypothesis, which has otherwise received poor to mixed support from previous studies that have mostly involved fewer species and shorter time series. Generalists achieve greater prevalence across the landscape, and we find some evidence for geographic variation in the abundance-diet breadth relationship, in particular among elevational bands. Interspecific variation in abundance also had a negative relationship with diet breadth, with specialists having more variable abundances across species. The fact that more specialized species can be both rare and common highlights the ecological complexity of specialization.

Keywords: Diet breadth, diet specialization, abundance, plant-herbivore, scale, tropics, long-term data, generalists, specialists

Introduction

      Insects are a remarkably diverse taxon, and all ecological guilds of insects engage in some of the most consequential and common ecological interactions. Ecology and evolutionary biology have benefited from studies of plant-insect interactions, especially studies on ecology and evolution of specialization, which have attracted decades of attention (Ehrlich and Raven 1964; Mani 1968; Forister et al. 2012). A common explanation and expectation for specialization has been that "a jack of all trades is a master of none," which is, in the ecological context of dietary specialization, the idea that a generalist might be able to consume a wide range of resources but will not be particularly well adapted to any one of them (MacArthur 1972; Futuyma and Moreno 1988). Two predictions are associated with this idea. First, functional trade-offs have long been hypothesized as the most parsimonious explanation for the predominance of specialists. If adaptation to one resource comes at the expense of lower performance on another resource, then specialists will be common and generalists will be favored only under certain circumstances. For example, a single-locus antagonistic pleiotropy for the ability to digest different food types favors specialization on one food type, whereas generalists will only be favored under special conditions, such as high fluctuations in host plant availability. That prediction has received only minimal support (Hardy et al. 2020), and is not our focus here. We are interested in a second prediction involving ecological consequences of specialization, specifically the idea that specialists should be locally more abundant, while generalists might never be as abundant in any one location but might occupy more places across broader landscapes. This prediction has often been stated in experimental terms for a small number of taxa (e.g., specialists will outperform generalists when tested on a common resource), which highlights the lack of relevant ecological, field-based datasets in the literature (Dyer et al.

2015). However, testing the empirical pattern in the field is of central importance to our

understanding of interaction networks and even ecosystem stability.

Studies that close the knowledge gap with respect to diet breadth and abundance at

different scales are likely to contribute substantially to our understanding of specialization and

the ecology and evolution of plant-insect interactions. Here, we use a 17-year empirical dataset

from Ecuador, which included replicated elevational gradients, to examine associations between

diet breadth and caterpillar (larvae of Lepidoptera) abundance, and to include potential effects of

spatial scale on these relationships. The long-term data include intensive sampling across space

and time, allowing for multiple measures of abundance. With these empirical data, we ask the

simple question of whether there is an association between abundance and diet breadth for

caterpillars in Ecuador and how the relationships between diet breadth and abundance vary

across different scales of observation. We hypothesized that caterpillar abundance would

decrease with increasing diet breadth, consistent with the "jack of all trades is a master of none"

hypothesis, especially at smaller spatial scales where the local demographic advantage of being

well-adapted to a host plant should be evident. At larger spatial scales, we expected to discover

an advantage to dietary generalization: because generalists can be found on multiple host species,

they should turn up in more surveys and plots, reflecting broader landscape-level occupancy.

Methods

Study location & collection protocol

We used 17-years of host plant-lepidopteran herbivore interactions collected within the

Eastern Andes of Ecuador between 2001–2018 (Fig. 1a; 420 km^2 from 00.36° S and 77.53° W)

(Wagner 2020) along an elevational gradient (100-3800m) that included low montane forests and

paramo habitat to explore relationships between caterpillar diet breadth and density. Caterpillars

were found by visual inspection of leaves using two distinct methods: general collecting and

sampling within standardized circular plots (10 m diameter). Both collection protocols and

rearing methods are fully described in previous publications (Dyer et al. 2007; Salcido et al.

2020). Briefly, general collection consisted of opportunistic collection of caterpillars along roads,

trails, transects, or off trail searches while plots were designated at least 10 m from roads or trails

and included estimates of plant diversity and genera-level leaf area. For both methods,

caterpillars were collected and assigned a specific record number and reared to adulthood.

Caterpillars and their host plant were identified to the lowest taxonomic level. In cases where the

resolution of taxonomic identifications was limited to family or genus, caterpillar or host plant

morphotypes were assigned based on family or genus designations in combination with character

descriptors. We removed data for which either host plant or caterpillar family and descriptors

were unavailable.

Diet breadth and density estimates

We used host plant and caterpillar taxonomic identifications from both general

collections and plots to calculate diet breadth of each caterpillar species. For our analyses, diet

breadth is the total number of unique host plants consumed by a given caterpillar species or

morphotype using all data from general collections and plots. We used only data collected within

plots to estimate caterpillar densities at plot and elevation scales. The abundance of each

caterpillar species was summed for each 10-m diameter plot (78.5 m^2). We also calculated an

index of landscape occupancy as the frequency of observation for a particular caterpillar species

across plots. Density per available host plant area and occupancy were summarized for plot and

elevation scales for which adequate data were available. Additionally, we explored an alternative

measure of density by calculating density per unit plot area and reported associated results in the

supplement.

Spatial scales

To evaluate the effect of spatial scale on associations between diet breadth and

abundance we examined patterns at various scales (in order of increasing area): 1) plot scale

(including plots that were aggregated to form larger subsets), 2) discrete elevational bands, and

3) a regional scale. For plots, data were collected from a total of 455 plots. We examined the

effect of scale on the relationship between diet breadth and density at the plot-level, by

aggregating plots into groups and calculating the average density and occupancy for each

caterpillar species across all plot groups. This process was iterated for plot groupings of

increasing size. To group them, a moving window method was applied (Fig. 1b). The window

size was equivalent to the number of plots grouped together (n=2-455 plots) and moved

sequentially (in order of time that the plots were completed) along the plot data in order to

maximize spatial and temporal similarities among plots. For each iteration, the window size

increased by one plot. For example, for a window size of two, caterpillar abundance and area

from plot 1 and plot 2 was summarized followed by all iterations of plot n-1 and plot n. In the

subsequent iteration, the window size increased to three plots and so on until the window size

equaled the number of all plots. Results presented in the main text are limited to groupings of 5

($\sim$393 m^2), 25 ($\sim$1963 m^2), 50 ($\sim$3927 m^2), and 250 ($\sim$19535 m^2) plots, however, results for all

possible aggregates were calculated and are presented in the supplementary text.

Plots within four distinct elevational bands: 1) 0 m - 1000 m, 2) 1000 m - 2000 m, 3)

2000 m - 3000 m, and 4) above 3000 m were aggregated (Fig. 1a), and these elevational bands

are ecologically very distinct (Pyrcz et al. 2009). Density was calculated based on the total

caterpillar abundance within each elevational band per available host plant area. While turnover

is substantive at a finer elevational scale (e.g., Sklenár and Ramsay 2001), elevational bands

were defined based on data availability and to allow for sufficient data at distinct elevations.

To examine caterpillar abundance at the regional level, total abundance for each

caterpillar species was summarized across the entire dataset. This included caterpillars collected

from both plot-based and opportunistic collection methods. With the data summarized at this

level, we needed another method to account for the fact that generalists could be observed more

often simply because they occur on a wider range of host plants. Thus, we calculated the number

of times that a caterpillar has been collected, divided by the number of hosts on which it occurs

(in other words, average abundance per host). For example, if a caterpillar A has a total

abundance of 10 and a diet breadth of 2 (it uses 2 host plant species), the density per host species

is 5. Prior to the analyses described here, we identified extreme outliers and singletons (species

for which there was only one record in the dataset) and removed them from statistical analyses.

Due to its unusually high abundance from sampling bias for related research projects, we also

removed *Eois olivacea* (Geometridae: Larentiinae).

Statistical inference

We used Bayesian linear models to estimate beta coefficients for the association between

abundance or occupancy and diet breadth for plot, elevational and regional scales. Models were

fit in JAGS (version 3.2.0) utilizing the rjags package in R (R version 3.6.3 and RStudio version

1.2.5042) using (for each analysis) two Markov chains and 1,000,000 steps each; performance

was assessed through examination of chain histories (burnin was not required), effective sample

sizes and the Gelman and Rubin convergence diagnostic. Response variables (abundance and

occupancy) were modeled with assumptions of normally distributed residuals with means

dependent on an intercept plus diet breadth as a predictor variable. We used weakly informative

priors: priors on beta coefficients were normal distributions with mean of zero and precision of

0.001 (variance = 1000); priors on precisions were modeled as gamma distributions with

rate = 0.1 and shape = 0.1. We report point estimates for beta coefficients and corresponding 95%

credible intervals (CI). Values for density, occupancy, and diet breadth were log transformed

($\log_{10}$) prior to analyses.

Results

Caterpillar density per available host area showed a negative relationship with diet

breadth for plot and elevational scales, except for the highest elevation band (Fig. 2a-b, Table 1,

Fig. S1). This relationship reflects a tendency for more specialized caterpillar species to be more

abundant wherever they are found. In contrast, occupancy (frequency of occurrence across the

landscape) showed a positive relationship with diet breadth for all plot cluster sizes (Fig. 2c,

Table 1, Fig. S2); however, the relationship varied across different elevation bands (Fig. 2d,

Table 1). A positive relationship was observed at the middle elevation bands (1000-2000 m and

2000-3000 m) while weak positive and negative relationships between occupancy and diet

breadth were observed at the low (< 1000 m) and high (>3000 m) elevation bands respectively

(Fig. 2d, Table 1). At our largest spatial scale, using all data, including plots and general

collections, we did not detect a relationship between density per available host plant species (β=

-0.076, 95% CI [-0.15, -0.0061] and diet breadth (Fig. 3). When density was calculated per plot

area, patterns at both plot and elevation scales were similar to trends observed for occupancy

(Fig. S3-4).

Discussion

As a theoretical expectation, the idea that a "jack of all trades is a master of none" has

withstood decades of examination with limited consensus, depending on the organisms and the

traits being examined (Futuyma and Moreno 1988; Forister et al. 2012). For many

morphological traits, biophysical constraints in vertebrates have produced clear examples where

an ecological generalist underperforms specialists (Oufiero et al. 2012). The story has been more

complicated for herbivorous insects, where the focal trait is typically developmental performance

in juvenile life history stages, and specialists have often not outperformed generalists when

experimentally evaluated on a common resource (García-Robledo and Horvitz 2012; Agrawal

2020). Instead of an experimental framework, we have taken a different approach by examining

the relationship between abundance and diet breadth measured across large spatial and temporal

scales in the field, and consistent with expectations we find that specialists tend to be locally

more abundant while generalists occupy more places on the landscape. This result raises the

possibility that specialists are indeed better adapted to their host plants, while generalists benefit

from being able to colonize more hosts. That general pattern is largely robust to observations

taken at different elevational and spatial scales (Fig. 2), although we did not detect a signal at the

very largest scale that included general collecting (Fig. 3), possibly because our measure of

"abundance" at that level is not as sensitive as the plot-level counts of individuals used at the

smaller spatial scales.

The other classic expectation from the jack-of-all-trades hypothesis is that the local

advantage of specialists derives from genetic trade-offs that prevent the maximization of fitness

across resources (Levins 1968; Agosta and Klemens 2009). That explanation has been

repeatedly questioned, with very few exceptions (e.g., Gompert and Messina 2016), and it is

important to note that the results reported here do not support or address the idea of genetic

trade-offs. Instead, it should be remembered that other explanations for a local specialist

advantage have been proposed (Fry 1996), including more efficient natural selection for

populations exposed to a single resource (Whitlock 1996). Regardless of the mechanism

explaining the tendency towards higher densities of more specialized species, an important

finding here is not only the sign and magnitude of that relationship, but also the shape of the

variance. In particular, the range of densities observed for specialists is orders of magnitude

greater than the range for densities of generalists (Fig. 2A). This broad distribution of densities

for specialists counterintuitively raises the possibility that being a host specialist is a more

multifaceted condition than host generalization. Specialists, for example, likely contend with a

different suite of natural enemies that affect local abundance in ways that are heterogenous and

deserve further study (Singer and Stireman 2005). Future investigations could, for example,

target dramatic differences that we report in the ratio of specialists to generalists that we report

among host genera (Fig. S5).

In conclusion, the results reported here are a reminder that many decades are often

needed for the evaluation of basic assumptions that underlie ecological theory, especially when

that theory is applied to complex traits such as digestion, development, and host use. A similar

study was recently conducted with armored scale insects, with the conclusion that diet breadth

and abundance are not linked (Peterson et al. 2020). Although Peterson et al. (2020) did not

detect an overall abundance-diet breadth relationship, it is the case that the most abundant

species tended to be extreme specialists. There is clearly much yet to be learned about ecological

and dietary specialization, and the observations reported here are made more relevant in the

midst of ongoing reports of declining insect abundances from around the world (Salcido et al.

2020; Wagner 2020). We hope that coming decades see long term datasets, such as the one used

here, being used simultaneously to motivate conservation and to test long-standing and

unresolved ideas about how the natural world works.

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Literature cited

Agosta SJ, Klemens JA. 2009. Resource specialization in a phytophagous insect: no evidence for genetically based performance trade-offs across hosts in the field or laboratory. J Evol Biol. 22(4):907–912. doi:10.1111/j.1420-9101.2009.01694.x.

Agrawal AA. 2020. A scale-dependent framework for trade-offs, syndromes, and specialization in organismal biology. Ecology. n/a(n/a):e02924. doi:10.1002/ecy.2924.

Dyer L, Massad T, Forister M. 2015. Dyer, L.A., T.J. Massad, and M.L. Forister. 2015. The question of scale in trophic ecology. Pages 288-317 in: Hanley, T. and K. La Pierre (eds.). Trophic Ecology: Bottom-Up and Top-Down Interactions across Aquatic and Terrestrial Systems. Cambridge University Press, Cambridge, MA.

Dyer LA, Singer MS, Lill JT, Stireman JO, Gentry GL, Marquis RJ, Ricklefs RE, Greeney HF, Wagner DL, Morais HC, et al. 2007. Host specificity of Lepidoptera in tropical and temperate forests. Nature. 448(7154):696–699. doi:10.1038/nature05884.

Ehrlich PR, Raven PH. 1964. Butterflies and plants: A Study in coevolution. Evolution. 18(4):586–608. doi:10.2307/2406212.

Forister ML, Dyer LA, Singer MS, Stireman JO, Lill JT. 2012. Revisiting the evolution of ecological specialization, with emphasis on insect–plant interactions. Ecology. 93(5):981–991. doi:10.1890/11-0650.1.

Fry JD. 1996. The evolution of host specialization: are trade-offs overrated? Am Nat. 148:S84–S107.

Futuyma DJ, Moreno G. 1988. The evolution of ecological specialization. Annu Rev Ecol Syst. 19(1):207–233. doi:10.1146/annurev.es.19.110188.001231.

García-Robledo C, Horvitz CC. 2012. Jack of all trades masters novel host plants: positive genetic correlations in specialist and generalist insect herbivores expanding their diets to novel hosts. J Evol Biol. 25(1):38–53. doi:https://doi.org/10.1111/j.1420-9101.2011.02401.x.

Gompert Z, Messina FJ. 2016. Genomic evidence that resource-based trade-offs limit host-range expansion in a seed beetle. Evolution. 70(6):1249–1264. doi:10.1111/evo.12933.

Hardy NB, Kaczvinsky C, Bird G, Normark BB. 2020. What We Don't Know About Diet-Breadth Evolution in Herbivorous Insects. Annu Rev Ecol Evol Syst. 51(1):103–122. doi:10.1146/annurev-ecolsys-011720-023322.

Levins R. 1968. Evolution in changing environments. New Jersey: Princeton University Press. [accessed 2020 Nov 20]. https://press.princeton.edu/books/paperback/9780691080628/evolution-in-changing-environments.

MacArthur RH. 1972. Geographical ecology: patterns in the distribution of species. New York: Harper & Row.

Mani MS. 1968. Ecology and biogeography of high altitude insects. Springer Netherlands (Series
{Entomologica}).

Oufiero CE, Holzman RA, Young FA, Wainwright PC. 2012. New insights from serranid fishes
on the role of trade-offs in suction-feeding diversification. J Exp Biol. 215(21):3845–3855.
doi:10.1242/jeb.074849.

Peterson DA, Hardy NB, Morse GE, Itioka T, Wei J, Normark BB. 2020. Nonadaptive host-use
specificity in tropical armored scale insects. Ecol Evol. doi:https://doi.org/10.1002/ece3.6867.
[accessed 2020 Nov 20]. https://onlinelibrary.wiley.com/doi/abs/10.1002/ece3.6867.

Pyrcz T, Wojtusiak J, Garlacz R. 2009. Diversity and Distribution Patterns of Pronophilina
Butterflies (Lepidoptera: Nymphalidae: Satyrinae) along an Altitudinal Transect in North-
Western Ecuador. Neotrop Entomol. 38:716-U3.

Salcido DM, Forister ML, Lopez HG, Dyer LA. 2020. Loss of dominant caterpillar genera in a
protected tropical forest. Sci Rep. 10(1):1–10. doi:10.1038/s41598-019-57226-9.

Singer MS, Stireman JO. 2005. The tri-trophic niche concept and adaptive radiation of
phytophagous insects. Ecol Lett. 8(12):1247–1255. doi:https://doi.org/10.1111/j.1461-
0248.2005.00835.x.

Sklenár P, Ramsay PM. 2001. Diversity of zonal páramo plant communities in Ecuador. Divers
Distrib. 7(3):113–124. doi:10.1046/j.1472-4642.2001.00101.x.

Wagner DL. 2020. Insect Declines in the Anthropocene. Annu Rev Entomol. 65(1):457–480.
doi:10.1146/annurev-ento-011019-025151.

Whitlock MC. 1996. The red queen beats the jack-of-all-trades: the limitations on the evolution
of phenotypic plasticity and niche breadth. Am Nat. 148:S65–S77.

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304 **Table 1.** Summary statistics of density per available host plant area and occupancy at plot and
305 elevation level.
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Level of observation	Density per available host plant area		Occupancy	
	β coefficient	95% CI	β coefficient	95% CI
Plot level (cluster size)				
5	-0.35	[-0.47, -0.22]	0.64	[0.56, 0.72]
25	-0.36	[-0.49, -0.23]	0.54	[0.46, 0.61]
50	-0.36	[-0.48, -0.23]	0.47	[0.39, 0.54]
250	-0.35	[-0.47, -0.24]	0.24	[0.15, 0.32]
Elevation level				
Low (<1000 m)	0.097	[-1.72, 1.63]	0.12	[-0.33, 0.50]
Mid 1 (1000-2000 m)	-0.31	[-0.49, -0.12]	0.43	[0.34, 0.53]
Mid 2 (2000-3000 m)	-0.35	[-0.48, -0.21]	0.62	[0.52, 0.71]
High (>3000 m)	-0.73	[-1.48, 0.068]	-0.049	[-0.30, 0.20]

Figure legends

Figure 1. This study took place in the Northeastern Andes in Ecuador, and plot-level data were

aggregated to examine the effect of increased scale on the relationship between density and diet

breadth. A) Samples were collected within the Napo Province across different elevations. Plots

were categorized into one of four distinct elevational bands (0-1000m, >1000-2000m, >2000-

3000m, >3000m from sea level); although substantial species turnover occurs at finer scales, we

only had sufficient data for these groups. Lines delineating the different elevations are not exact

and are intended to be used as a heuristic tool. B) Effects of scale were evaluated at the plot level

by sequentially aggregating plots (circles) using a moving window method (solid and dashed

boxes). The moving window method was applied to the plot data iteratively: each time the size

of the window (the number of plots aggregated) increased by one more plot. For a window size

of five (e.g., plots 1-5), every two plots were sequentially aggregated. In the next iteration, every

six plots were sequentially aggregated (e.g., plots 1-6). For each aggregate and caterpillar species

(circular nodes), abundance was summarized. Density was calculated based on the combined

area or corrected for the leaf area of associated host plant genera (square nodes). Density across

all plot aggregates for a given species were averaged.

Figure 2. Relationship among diet breadth and caterpillar density corrected for available host

area (cm^2) and occupancy (landscape presence) at different scales of observation: a) and c) are

plot scales, and b) and d) are elevation scales. Plot-level data were combined for each caterpillar

species (points) for various plot aggregates (blue: 5 plots, red: 25, green:50 and purple:250) to

evaluate the effect of scale. For the elevation analysis, discrete elevation bands included: 0 -

1000 m (red), 1000-2000 m (green), 2000-3000 m (purple), and above 3000 m (blue). Points

were set to a transparency value given the quantity of points, such that darker hues indicate a

high density of points.

Figure 3. Relationship between caterpillar density per host plant species and diet breadth at a

regional scale. Points were set to a transparency value given the quantity of points, such that

darker hues indicate a high density of points.

367   Figure 1

376   Figure 2

378   Figure 3

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