

Adult and larval traits as determinants of geographic range size among tropical reef fishes

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Most marine organisms disperse via ocean currents as larvae, so it is often assumed that larval-stage duration is the primary determinant of geographic range size. However, empirical tests of this relationship have yielded mixed results, and alternative hypotheses have rarely been considered. Here we assess the relative influence of adult and larval-traits on geographic range size using a global dataset encompassing 590 species of tropical reef fishes in 47 families, the largest compilation of such data to date for any marine group. We analyze this database using linear mixed-effect models to control for phylogeny and geographical limits on range size. Our analysis indicates that three adult traits likely to affect the capacity of new colonizers to survive and establish reproductive populations (body size, schooling behavior, and nocturnal activity) are equal or better predictors of geographic range size than pelagic larval duration. We conclude that adult life-history traits that affect the postdispersal persistence of new populations are primary determinants of successful range extension and, consequently, of geographic range size among tropical reef fishes.

macroecology | marine dispersal | colonization

Geographic range size is a fundamental biogeographic variable that, among other effects (1, 2), strongly influences a species susceptibility to extinction (3, 4). Because most marine organisms disperse as larval propagules transported by ocean currents, it is often assumed that the duration of the larval stage is the fundamental determinant of their dispersal ability, and hence their range size (5, 6). Tropical reef fishes have geographic ranges that vary greatly in size, from a few square kilometers around tiny isolated islands to entire ocean basins (7–9). Given that pelagic larval duration (PLD) also varies greatly among such fishes, from only a few days to many months, the effects of PLD on dispersal potential became an early focus of investigation on general determinants of range size among those fishes and other near-shore marine species (10–12). However, although it has become evident that PLD is unlikely to be a primary determinant of geographic range size (13–16), alternative hypotheses have only recently begun to be considered (9).

To expand its geographic range, a species must successfully colonize new areas following the dispersal of its propagules (17). Consequently, attributes other than pelagic dispersal capacity may largely determine how widely reef fishes are distributed geographically (9). Here we assess the relative importance of seven adult and larval traits in influencing geographic range sizes of tropical reef fishes at the global scale. We do so using data from 590 species of tropical reef fishes in 47 families, the largest compilation of such data currently available for any marine group (Dataset S1). Traits directly linked to larval dispersal potential include PLD and spawning mode. Adult traits include maximum body size, schooling behavior, nocturnal activity, use of multiple habitat types, and adult depth range. The adult-biology traits chosen are not directly related to larval dispersal

potential, but may influence the propensity for range expansion by affecting the establishment and persistence of new populations, as suggested by a recent study on Atlantic reef fishes (9). For example, schooling (18, 19) and nocturnal activity (20) reduce predation risk and thereby increase the chance of post-settlement survival. Broad habitat use and depth range indicate ecological generality, which is thought to influence establishment success in new environments (21). Finally, body size is linked to both predation risk and ecological generality (22).

Evaluation of these hypotheses is challenging because species traits are phylogenetically nonindependent (23) and unevenly distributed among families. Previous studies of dispersal-range-size relationships have controlled for effects of phylogeny, and limits on range-size imposed by ocean-basin size, by separately analyzing subsets of data (7, 16). However, this approach reduces statistical power (23, 24) and the ability to assess the generality of the effects of different factors. Our analysis controls for the nonindependence of shared traits among related species by using linear mixed-effects modeling (LMM) treating family and genus as nested random effects (9, 23). Our analysis includes species from three different regions that vary greatly in maximum (longitudinal or latitudinal) extent: the Indo-Central Pacific (ICP; ~22,000 km), the tropical Atlantic (TA; ~12,000 km), and the tropical eastern Pacific (TEP; ~5,000 km). To control for this variation, we include region and its interactions with other variables as fixed effects in our models. Modeling the data in this way, we are unique in being able to assess the relative importance of various adult and larval traits as determinants of range size among tropical reef-fish, as a group, at the global scale.

Significance

Marine organisms disperse mostly by ocean currents as larval propagules. Therefore, it is commonly thought that the duration of the larval stage is the fundamental determinant of geographic range size. Using a global compilation of reef fish traits, we test an alternative hypothesis: adult traits associated with population establishment and persistence in novel areas are better predictors of geographic range size than larval traits. We conclude that colonization success is as primary determinant of successful range extension and of geographic range size among tropical reef fishes.

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Results and Discussion

Our analysis shows that region is the most significant predictor of geographic range size (Table 1). This result could be because of constraints of regional extent on range size within a region, or to differences in the underlying structure of the range-size frequency distribution within each region. To assess the importance of regional extent, we repeated the analysis after expressing the range size of each species as the ratio between the range size of species and the extent of the region. When range sizes were standardized in this way, all effects of species traits were maintained, and the region effect declined from the most important to the least important predictor (Table 1).

Our analysis also shows that three of five adult life-history traits—body size, schooling behavior, and nocturnal activity—are significant predictors of geographic range size (Figs. 1 and 2) for the global dataset (Table 2). Although PLD is also a significant predictor of range size at the global scale, only the largest of the three regions, the ICP, showed such a correlation (Table 2 and Table S1), confirming the finding of a previous analysis (7). The ICP correlation is largely driven by the few transpacific species that cross the world’s largest oceanic barrier, the 4,000+-km-wide Eastern Pacific Barrier (EPB) that separates the ICP from the TEP (7). After the removal of those trans-Pacific species, the effect of PLD dropped from the second-most influential factor, globally, to the least-important trait (Table 1). A recent analysis of global ocean circulation patterns and habitat distributions (16) indicates that larvae of most tropical-reef species have PLDs sufficient to reach most habitat patches. Our results are consistent with that analysis: they show that the PLD effect is evident only at the largest spatial scale and when habitat patches are most isolated.

Our analysis provides further evidence for the view that range extensions are strongly influenced by adult life-history traits (9), factors likely to affect the capacity of new colonizers to survive and establish reproductive populations. All three of the positive adult-biology correlates of range size we identified—maximum body size, schooling behavior, and nocturnal activity—may enhance the probability of population establishment after propagule arrival.

Predation is one of the main processes influencing the structure and species composition of ecological communities (25), especially assemblages of coral reef fishes (26). If these factors limit the number of locally coexisting species (27, 28), predators may inhibit nonnative species establishment (29), and thereby

constrain the geographic ranges of prey (30). Piscivorous predators are ubiquitous in coral reefs. Mortality is disproportionately higher among new recruits (31, 32), although juveniles and adults are not immune to predation, as evidenced by morphological, chemical and behavioral antipredatory mechanisms they use (26, 33). Among reef fishes, predation may be particularly important for species at early-stage colonization if predators tend to target rarer species (27, 34). Immigrant species may have better chances of survivorship and establishment if they can rely on specific antipredator mechanisms.

Reduction in predation risk is considered one of the main benefits of schooling in fish (35), and per capita mortality rates in schools of reef fishes decrease as school size increases (36). Predators are less successful at singling out individual prey from large schools because of the “confusion effect” (34). Survivorship may increase in both single-species schools and mixed-species schools not only through the confusion effect but also through more effective foraging by social observation, better vigilance for predators, and greater economy of time budgeting (35). Mixed-species schools are common among recruiting juveniles of reef fishes, sometimes involving species with different diets, which reduces the competitive costs while retaining the benefits of social behavior (19). Schooling may also increase the chance of finding mates among a scarce cohort of colonizers, improve food detection, and enhance access to resources protected by territorial competitors (37, 38). Thus, schooling fishes may have greater relative potential for population establishment and persistence after reaching new areas.

Recently, direct evidence has emerged that diel activity patterns of reef fishes are influenced by ongoing predation intensity: day-active nocturnal fish were much more common at a predator-depleted atoll in the Central Pacific than at a neighboring atoll with a large population of predators (20). This finding supports the view that nocturnal activity allows prey to avoid interactions with day-active predators. Differences in the morphology and behavior of nocturnal and diurnal reef fishes also are consistent with reduced predation intensity at night (39). Nocturnal planktivores are relatively deep bodied and robust, in contrast to their more streamlined diurnal counterparts, and there is a general reduction of schooling at night (39). Whatever the ultimate reasons for the development of nocturnal activity among reef fishes, nocturnal species may be exposed to a smaller subset of predators than day-active fishes (26, 39).

Table 1. Significant variables ranked according to their independent effects

All species (n = 590)					Trans-Pacific species removed (n = 564)				
Variable	df	F value	P value	IE (%)	Variable	df	F value	P value	IE (%)
Geographic range size									
Region	2	117.83	<0.001	53.3	Region	2	104.80	<0.001	66.9
PLD	1	54.72	<0.001	17.3	Body size	1	31.44	<0.001	9.6
Body size	1	69.86	<0.001	15.5	Nocturnal	1	16.69	<0.001	8.4
Nocturnal	1	18.22	<0.001	7.8	Schooling	1	15.45	<0.001	8.0
Schooling	1	12.63	<0.001	6.0	PLD	1	12.23	<0.001	6.9
Ratio between the species range size and region size									
Body size	1	111.70	<0.001	37.6	Body size	1	90.06	<0.001	36.0
PLD	1	23.28	<0.001	24.8	Nocturnal	1	38.13	<0.001	25.9
Nocturnal	1	33.17	<0.001	19.4	Schooling	1	12.94	<0.001	16.2
Schooling	1	11.47	<0.001	13.9	PLD	1	11.62	0.001	13.3
Region	2	2.09	0.123	4.2	Region	2	3.48	0.031	8.3

The independent effects (IE) value corresponds to the percentage of the explained variance accounted for by each explanatory variable as calculated using hierarchical partitioning. In Geographic range size, ranking is according to IE on geographic range size. In the ratio between the species range size and region size ranking is according to IE on the ratio between the species range size and region size. Degrees of freedom (df), test statistics (F value), and probabilities (P value) are listed for each coefficient in each model.

Table 2. Parameters of the final predictive LMM

Variable	Estimate	SE	t value	P value
Intercept	987.76	4425.75	0.223	0.823
Body size	2210.34	335.41	6.589	<0.001
Schooling behavior	2030.28	609.71	3.329	0.001
Nocturnal activity	2503.27	881.27	2.840	0.004
PLD: Tropical Atlantic	-247.54	1295.63	-0.191	0.848
PLD: Indo-Central Pacific	4935.42	1434.00	3.441	<0.001
PLD: Tropical Eastern Pacific	501.14	2406.30	0.208	0.835
Region: Indo-Central Pacific	-10483.26	4915.94	-2.132	0.033
Region: Tropical Eastern Pacific	-5102.71	8397.86	-0.607	0.543

Parameters estimated in a LMM with MLD of geographical range size (in kilometers) as the response variable for larval and adult traits including an interaction between PLD and biogeographic region, and genus nested within family as a random variable. Estimate, coefficient estimate of explanatory variables; SE, test statistic (t value) and probability (P value). P values are significant ($P < 0.05$) for coefficients if in bold. Reference levels for this regression were set as "Tropical Atlantic" for region and as "no" for both schooling behavior and nocturnal activity.

species in response to ocean warming have been based on their thermo-physiological tolerances (49). However, our results indicate other life-history traits that may constrain range extension and thereby influence extinction risks (4).

Materials and Methods

Species Traits. We selected life-history traits that are thought to potentially influence range sizes (2, 9), and for which data are readily available for all species for which we also have PLD data. Body size is correlated with many other biological attributes of species, including range size (2). Schooling behavior and nocturnal activity are mechanisms that may diminish predation risk (18–20), and thereby enhance range expansion by assisting colonization. Environmental generalist species might be expected to have greater establishment success than specialists in new and different environments (21). Here, such generalists included species that have larger depth ranges and those that use other habitats in addition to structural reefs (soft bottoms, seagrass/macroalgae beds, mangroves, and estuaries).

PLD is linked to the time larvae are susceptible to pelagic dispersal. Spawning mode (production of pelagic or benthic eggs), besides acting as a proxy for PLD (9), influences the stage of development at which larvae enter the pelagic zone. The eggs of pelagic spawners are immediately subjected to transport by currents, whereas the eggs of benthic spawners are deposited on the substrate and the larvae enter the pelagic realm only after developing for some time in the egg before hatching. Existing data on tropical reef fish PLD were compiled from the primary literature for 446 species (see references in [Dataset S1](#)). Additional data on PLDs were obtained for 227 species, from which 144 are uniquely reported, by aging settlers through analyses of daily growth increments and settlement marks in otoliths (50). Because of a lack of sufficient data in one or more regions we did not include one other factor that we assessed in our study of range expansion across large barriers in the Atlantic Ocean (9): flotsam-rafting behavior by postlarval stages. Data on spawning mode, maximum total length (our metric for body size), depth range, schooling behavior, multi-habitat use, and nocturnal activity were obtained from the primary literature and the global fish data aggregator FishBase (www.fishbase.org). We considered schooling species to be those that regularly form polarized, cohesive groups of 20 or more individuals. Diurnal and nocturnal were defined by the main period of day that each species actively forages.

Geographic Range. For comparability with previous studies we used the linear distance between the farthest two range endpoints—maximum linear distance (MLD) in kilometers—as a metric of geographic range size (7, 16). Those studies found that the MLD of range size is strongly correlated with the combined maximum latitudinal and longitudinal extent of ranges, and they considered MLD to represent an adequate descriptor of a species' geographic extent (7). Data on species range limits were obtained from guidebooks (51–55) and from the Ocean Biogeographic Information System (OBIS), a global aggregator of geo-referenced collection records (www.iobis.org). Data from both FishBase and OBIS were screened by us and complemented from our own records (56) ([Dataset S1](#)). Endpoint geographic coordinates were determined to the nearest degree using Google Earth (earth.google.com). The MLD was measured using the function "geodist" in the R package "gmt" (57). Species were grouped in terms of their residence in three well-defined biogeographic regions, the physical dimensions of which delineate the maximum attainable range size in each region: the Indo-central Pacific region (from the Red Sea to Easter Island), the tropical Atlantic (from the northwest Gulf of Mexico to the southern Gulf of Guinea), and the tropical eastern Pacific (from the northern Gulf of California to northern Peru). Primarily tropical species that extend their ranges to temperate zones had their full range considered in the analysis. Although it is in the Pacific, together with the ICP, the TEP is a well-recognized biogeographic entity that has a substantial fauna with a high level of endemism (~75%) (47). Only TEP endemics were included in the TEP group in our analyses, which avoided possible confounding effects of recent trans-Pacific crossers not having realized their full range potential in that region. The few trans-Pacific species that occur in both the ICP and TEP were included as members of the ICP fauna because most of the range of each is in the ICP and most appear to have migrated from there to the TEP (48). We did not consider the western and eastern sides of the TA as separate regions because there are insufficient data on the PLDs of species endemic to the tropical eastern Atlantic, which has a substantially lower rate of endemism than the tropical western Atlantic (58).

Statistical Analysis. The LMM was fitted using the function "lme" from the package "nlme" (59) in R. The response variable was the MLD between two points along the geographic range boundary. PLD, spawning mode, maximum body size, schooling behavior, nocturnal activity, depth range, multi-habitat use, and region were included as fixed variables, and genus and family were included as nested random variables. Interactions between region and each fixed factor were considered in the full model. For model selection, we followed the procedure recommended by Zuur et al. (60) of a backward stepwise removal of nonsignificant fixed-effect terms ($P > 0.05$) from the full model based on log-likelihood ratio tests ([Table S1](#)). Partitioning of variance to determine the relative importance (percent of explained variance; independent effects in [Table 1](#)) accounted for by each explanatory variable in the model was calculated by hierarchical partitioning using the R package "hier.part" (61).

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