

THE EFFECT OF INVASIVE LIONFISH ON REEF FISH COMMUNITY STRUCTURE
ALONG THE MESOAMERICAN BARRIER REEF

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ABSTRACT

Serena Hackerott: The effect of invasive lionfish on reef fish community structure along the Mesoamerican Barrier Reef
(Under the direction of John F. Bruno)

Lionfish are invasive predators, native to the Indo-Pacific, assumed to be negatively affecting Caribbean coral reefs. Small-scale studies suggest lionfish can reduce the abundance and diversity of small prey individuals. However, it is unclear whether lionfish predation affects entire reef fish communities. Our goal was to assess the effect of lionfish on coral reef fish communities across a complex reef system. We quantified fish abundance, diversity, and community composition at sixteen reefs along ~250km of the Mesoamerican Barrier Reef in Belize over five years, including the onset of the invasion. Lionfish had no effect on reef fish community structure on our sites in Belize. The effects of lionfish may be density dependent, and current densities in Belize are likely due to a combination of natural factors and lionfish removals. Because current densities have no effect on reef fish communities, additional lionfish removal efforts may not be necessary to prevent impacts.

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LIST OF ABBREVIATIONS

AIC	Akaike's "An Information Criterion"
ANOSIM	Analysis of Similarity
BACI	Before After Control Impact
MBR	Mesoamerican Barrier Reef
MPA	Marine Protected Area
NMDS	Non-metric Multidimensional Scaling
NTZ	No-Take Zone
PERMANOVA	Permutational Multivariate Analysis of Variance
SE	Standard Error
TL	Total Length
VIF	Variance Inflation Factor

CHAPTER 1: THE EFFECT OF INVASIVE LIONFISH ON REEF FISH COMMUNITY STRUCTURE ALONG THE MESOAMERICAN BARRIER REEF

Introduction

Invasive species are widely recognized as a major threat to biodiversity and an important conservation priority (Bax et al. 2003, Pimentel et al. 2005, Molnar et al. 2008, Vilà et al. 2011). While the qualities conferring invasiveness of a species or “invasibility” of an ecosystem have been thoroughly explored, mostly in terrestrial ecosystems (Rejmánek and Richardson 1996, Levine and D’antonio 1999, Lonsdale 1999, Richardson and Pyšek 2006, Van Kleunen et al. 2010), studies quantifying the factors influencing the impact of an invasive species on the recipient community are less represented (Parker et al. 1999). Furthermore, the invasiveness of a species does not predict the impact it will have on invaded ecosystems (Ricciardi and Cohen 2007). Studies of invasion success can contribute to policies aimed at preventing invasions, but impact-focused studies are essential to developing conservation strategies once an invasive species has already become established.

The impact of an invasion can also vary between invaded habitats, specifically across gradients of habitat connectivity. When exotic predators become invasive within isolated systems, prey populations are often decimated. For example, feral cats introduced to islands have caused an estimated 14% of the global extinctions of birds, mammals, and reptiles (Medina et al. 2011). However, invasive predators may have less of an impact on prey communities within more connected, open systems due to re-colonization through meta-population dynamics or “the rescue effect”(Gotelli 1991).

Indo-Pacific lionfish (*Pterois volitans* and *Pterois miles*, hereafter called “lionfish”) are one of the first exotic predators to invade an open marine system, and the first to become established across the Caribbean region (Schofield 2009). Several characteristics such as cryptic coloration (Kindinger 2014), undetectable chemical cues (Lönngstedt and McCormick 2013), and novel predation tactics (Albins and Lyons 2012), may increase their effectiveness as predators in their invaded range. Due to their generalist diet, consuming over 100 Caribbean species (Morris and Akins 2009, Green et al. 2012, Green and Côté 2014), and high consumption rates (Cote and Maljković 2010), lionfish have the capacity to affect numerous native species and drive changes in native fish communities. In fact, lionfish significantly decreased the abundance, species richness, and diversity of coral reef fish recruits (<5 cm TL) during eight weeks on experimental reefs off of Lee Stocking Island, Bahamas (Albins 2013). Additionally, increasing lionfish abundance has been associated with a significant decrease in the biomass of native prey individuals (<15 cm TL) over two years along a continuous reef in New Providence, Bahamas (Green et al. 2012). However, these and other previous studies have only demonstrated the negative effects (i.e., decline in abundance and diversity) of invasive lionfish on small, native prey individuals on relatively small, isolated systems and controlled experimental reefs (Albins and Hixon 2008, Green et al. 2012, Albins 2013). It has yet to be determined whether the predation pressure of lionfish on small prey individuals drives changes in the entire (i.e., all size classes) reef fish community.

The goal of our study was to quantify the effect of the lionfish invasion on reef fish communities across the world’s second largest barrier reef system: the Mesoamerican Barrier Reef (MBR). We tested whether reported effects of lionfish on small native fishes at small

spatial scales (Albins and Hixon 2008, Green et al. 2012, Albins 2013) can be detected on the entire reef fish community on contiguous reef habitats, within the context of existing variability. Specifically, we investigated the effects of lionfish abundance on reef fish abundance, diversity, and community composition. Our 16 fore-reef sites were along the Mesoamerican Barrier Reef in Belize, spanning ~250 km, across gradients of natural environmental conditions as well as anthropogenic stressors. Our longitudinal study began in 2009, included the onset of the lionfish invasion in Belize (Schofield 2009), and continued until 2013. The design is a quasi-BACI design (Before After Control Impact) as we tracked changes in native fishes pre- and post-invasion and among sites varying in lionfish density over time. We hypothesized that reef fish abundance, species richness, and diversity would be negatively related with lionfish abundance due to the predation of invasive lionfish on small prey individuals (Albins and Hixon 2008, Green et al. 2012, Albins 2013). We also hypothesized that coral reef fish communities would experience a shift in composition due to the invasion as lionfish can differentially reduce the biomass of prey species (Green et al. 2012) by selectively feeding on species with certain morphological and behavioral traits (Green and Côté 2014).

Lionfish can potentially influence reef fish communities both directly through predation and indirectly through competition. Lionfish may also influence native fish communities by other indirect mechanisms such as acting as a disturbance that increases the availability of resources or niches for the remaining reef fish species (Ward and Stanford 1983). Therefore, to assess the effect of lionfish on native communities, we tested our hypotheses on two reef fish community groups. We first tested our hypotheses in terms of the reef fish community as a whole to quantify the overall (i.e. both direct and indirect) effect of

lionfish on reef fish. We then focused on the known prey fish community, or fish species previously documented as lionfish prey, to determine the direct effect of lionfish on reef fish communities.

Methods

2.1 Study Sites and Reef Fish Surveys

We monitored 16 fore-reef sites at 12 to 15 m depth along the MBR in Belize (Fig. 1, Table S1). Three sites were in No-Take-Zones (NTZ's) where no fishing is permitted and five sites were in Marine Protected Areas (MPA's) where limited fishing is permitted. Sites in either NTZ's or MPA's are hereafter referred to collectively as "protected" sites. Eight sites were not under fishing or use restrictions and are hereafter referred to as "unprotected" sites. To minimize habitat variability of survey sites, we only surveyed spur-and-groove habitats formerly dominated by *Orbicella* (formerly *Monstastrea*). At each site, underwater visual surveys of reef fish were performed using six to eight belt transects (modified from AGRAA v5.0 (Lang et al. 2010)) placed parallel to the spur-and-groove formations. We counted and identified reef fish to species, and estimated sizes (total length (TL)) in 5-10 cm intervals. Fish <5 cm TL, were counted in 15 x 1 m belt transects while fish between 5 and 40 cm TL were counted in 30 x 2 m belt transects. Fish >40 cm TL as well as lionfish were counted in 50 x 10 m belt transects, including a thorough search for cryptic lionfish. At each site, the longer and wider transect contained the smaller transects. We surveyed our sites each year from 2009-2013 in mid-May to early June during full daylight. Surveys were performed by the same team in 2010, 2012, and 2013 so only this data was included in our models (detailed below) to minimize potential variability in data collection due to surveyor bias.

2.2 Reef Fish Community Structure

Reef fish abundance (individuals/m²) was calculated for each reef fish species and total reef fish abundance was defined as the sum of the abundances of each species (excluding lionfish) per transect. Species richness and species diversity of reef fish were calculated to quantify changes in alpha diversity. Species richness (S) was calculated as the total number of species present (excluding lionfish) per transect. Species diversity was calculated in terms of Shannon-Wiener diversity (H'), a metric that accounts for both species richness and evenness, was calculated as:

$$- \sum_{i=1}^S p_i \cdot \ln(p_i)$$

where S = species richness, and p_i = the proportion of species i in the transect.

Community composition was quantified in terms of Bray-Curtis dissimilarity as:

$$\frac{2C_{ij}}{S_i + S_j}$$

where C_{ij} = species in common between sites i and j , and $S_{(i \text{ or } j)}$ = species in site i or j .

All indices of community structure and composition were quantified using the *vegan* (v2.0-10) package in R (R Development Core Team 2013).

2.3 Prey Community Structure

Prey species were identified as species documented to be consumed or directly influenced by lionfish throughout the Caribbean (Table S2) (Albins and Hixon 2008, Morris and Akins 2009, Layman and Allgeier 2011, Green et al. 2012, Valdez-Moreno et al. 2012,

Albins 2013, Cote et al. 2013, Green and Côté 2014). Prey fish abundance, species richness, species diversity, and community composition were quantified using the same methods previously described for the total fish community.

2.4 Abiotic Covariates

To account for variability among sites, we included three site-specific abiotic covariates in our analysis: site protection status, reef complexity, and humans per area of reef. All three are known to affect reef fish diversity and abundance (Friedlander et al. 2003, Williams et al. 2008) and could influence the effect of lionfish on fish communities. Due to regulations on fishing, protected sites may have higher fish abundance, species richness, or diversity than unprotected sites. Yet lionfish could have a weaker effect on reef fish communities on protected sites compared to unprotected due to lower abundances (Green et al. in press, Barbour et al. 2011, Frazer et al. 2012, Hackerott et al. 2013) or changes in lionfish behavior (Côté et al. 2014) due to lionfish culling. We split protected sites into either NTZ or MPA status as different protection levels may differ in influence on reef fish communities and the effect of lionfish. We included reef complexity as a predictor because highly complex reefs may support higher reef fish abundances or diversity compared with less complex reefs (Friedlander et al. 2003). Although reef complexity may not influence lionfish abundance (Valdivia et al. 2014, Anton et al. 2014), it could alter predation success of lionfish on resident reef fish by decreasing predation risk due to higher refuge availability for prey (Beukers and Jones 1998). To estimate reef structural complexity, we used an index from 0 to 5, where “0” was a reef with no vertical relief and “5” was an exceptionally complex reef (Polunin and Roberts 1993, Valdivia et al. 2014). Reef complexity was estimated along each transect and then averaged to obtain a single value for each site. Human

population per reef area (hereafter referred to as “humans/reef”) was used as a proxy for fishing pressure (Newton et al. 2007, Mora 2008). We hypothesized that fishing pressure could negatively affect reef fish abundance, species richness, and diversity, but would likely not influence lionfish abundance (Valdivia et al. 2014). Humans/reef was calculated as the human population within a 50 km radius of each site, divided by reef area within a 10 km radius of each site (details in Valdivia et al. 2014). A Spearman correlation matrix among the numerical explanatory variables indicated no correlation that could compromise simultaneous modeling, therefore we included all three abiotic covariates in the analytical models.

2.5 Analysis of the Effects of the Lionfish Invasion

We used generalized linear mixed effect models (glmer) to evaluate the effect of lionfish abundance and time since invasion on reef fish (total fish and prey) abundance, species richness, and species diversity, accounting for site-specific abiotic covariates. All reef fish community responses were modeled separately using the *lme4* (v1.1-5) package (R Development Core Team 2013). Total fish and prey fish abundance were modeled with a gamma log-link distribution, species richness with a gamma identity-link distribution, and species diversity with a Gaussian log-link distribution. Outliers, points outside of the “whiskers” in a boxplot of the values (split by year), were removed from lionfish abundance data (~4%, 17 total transects out of the 406 collected from 2009-2013) in all models to aid with model fit (Fig. S1). Additionally, the outliers of total fish (~7.5%, 16 transects out of the 214 collected in 2010, 2012, and 2013 and included in the models) and prey fish (~7%, 15 out of 214) abundance as well as diversity (~1%, 2 out of 214, for total fish and ~2%, 4 out of 214, for prey fish) were also removed from each respective model to aid with model fit

(Fig. S1). Site was included in each model as a random effect to account for within site variability. The number of years following the lionfish invasion, lionfish abundance, and the three abiotic covariates were coded as fixed effects in each model. We standardized (centered and divided by standard deviation) all numerical predictors in each model. We also included interactions between lionfish abundance, year, and each abiotic covariate in each global model. Global models were structured as:

$$\begin{aligned}
 y = & \beta_0 + \beta_1 \cdot Year_i + \beta_2 \cdot Lionfish\ Abundance_i + \beta_3 \cdot Protection\ Status_i \\
 & + \beta_4 \cdot Reef\ Complexity_i + \beta_5 \cdot \frac{Humans}{Reef}_i + \beta_6 \cdot Lionfish\ Abundance_i : Year_i \\
 & + \beta_7 \cdot Lionfish\ Abundance_i : Protection\ Status_i + \beta_8 \\
 & \quad \cdot Lionfish\ Abundance_i : Reef\ Complexity_i \\
 & + \beta_9 \cdot Lionfish\ Abundance_i : \frac{Humans}{Reef}_i + \varepsilon_i \approx N(0, \sigma_i^2)
 \end{aligned}$$

where $y = \log(u_i)$ for abundance and species diversity and $y = u_i$ for species richness of all fish and prey fish.

Each global model was dredged using the *MuMIn* (v1.9.13) package in R (R Development Core Team 2013) to extract all potential combinations of explanatory variables that produce top models with delta AIC < 2 (Burnham and Anderson 2002). Based on these selected models, coefficient estimates were averaged for each explanatory variable. The residuals of each final model were plotted against the fitted values and each of the predictors to check for homogeneity of variance of the errors. Variance Inflation Factor (VIF) were calculated for each predictor and threshold of VIF < 2 (Graham 2003) was used to check for

colinearity among the covariates included in final each model. Quantile-quantile plots (*car package v2.0-19*) were used to check normality of the residuals of each final model. Spline-correlograms (*ncf package v1.1-5*) using the location coordinates of each site with 1,000 resamples revealed no evidence of spatial autocorrelation of the residuals of each final model (R Development Core Team 2013).

Changes in reef fish (both total and prey fish) community composition in response to year (pre-invasion: 2010 and post-invasion: 2013) and lionfish abundance were first assessed using permutational multivariate analysis of variance (PERMANOVA) of the formula

$$[Transect \text{ by Species Abundance Matrix}] \sim Year_i * Lionfish \text{ Abundance}_i$$

with 10,000 permutations, stratified by site i , using Bray-Curtis dissimilarity (*vegan package v2.0-10*). Pre-invasion (2010) and post-invasion (2013) community compositions across all sites were quantified using an Analysis of Similarities (ANOSIM) with 10,000 permutations, grouped by year, using Bray-Curtis dissimilarity (*vegan package v2.0-10*). Reef fish community composition across all sites, pre- and post-invasion, were then assessed visually in two-dimensional space using nonmetric multidimensional scaling (NMDS) ordination analysis of Bray-Curtis dissimilarity using the *vegan (v2.0-10)* package in R (R Development Core Team 2013). We used NMDS because this ordination technique does not assume linearity of species responses or that species responses are due to environmental gradients (Minchin 1987).

Results

Among the three survey years included in the analysis, 128 species of reef fish were identified, 36 (28%) of which were documented as lionfish prey species (Table S2). No

lionfish were recorded on our sites in 2009 but by 2010, lionfish were present on two sites (Calabash and South Middle Caye) at relatively low abundances (6.7 ± 6.7 individuals/ha (mean \pm SE) on both sites), marking the onset of invasion. By 2012 lionfish were observed on 14 of 16 sites and on all sites by 2013. Across all sites, the average lionfish abundance (mean \pm SE) was 0.9 ± 0.7 individuals/ha in 2010, peaked to 21.27 ± 5.50 individuals/ha in 2011 and decreased slightly, but not significantly, in 2012 to 16.3 ± 5 individuals/ha and 11.1 ± 4.2 individuals/ha in 2013 (Fig. 2). The highest lionfish abundance found during our study was 70 ± 29.2 individuals/ha at Pampion in 2012. Due to the presence of lionfish on all 16 sites by the end of the study, our design was not a BACI design in a strict sense. However, lionfish density varied among sites over time and we used this variability to test the effects of lionfish over a gradient of potential impact levels (assuming the effects of lionfish are density-dependent).

Lionfish abundance had no significant effect on the abundance, species richness, or species diversity of total fish or prey fish communities (Fig. 3, Table S3). Reef complexity was positively related to total fish abundance and species richness (Fig. 3, Table S3). For the analysis of prey fish species richness, the null model was retained during the model averaging procedure which indicates that prey fish species richness is more strongly influenced by factors other than those included in our model. Total fish species diversity decreased (marginally significantly) with years following the lionfish invasion but this trend was not related to lionfish abundance. Total fish species diversity was also significantly lower on sites within MPA's or NTZ's compared to unprotected sites (Fig. 3, Table S3). The quantile-quantile plot of the residuals of the model of total fish species diversity indicated

that the residuals were slightly non-normal due to a few values at the extremes (Fig. S3), yet this was the best fit model for our available data.

The PERMANOVA analysis indicated total fish and prey fish community composition changed over time on some sites (p-value= $10.0e^{-5}$ and $10.0e^{-5}$, respectively) but community composition was not related to lionfish abundance (p-value= 0.87 and 0.83, respectively). Additionally, there were no clear patterns indicating changes in total fish or prey community composition from before (2010) to after (2013) the lionfish invasion across all sites (Fig. 4). This was confirmed by an analysis of similarity (ANOSIM) comparing communities across all sites between pre- and post-invasion for both total and prey fish (p-value= 0.61 and 0.53, respectively).

Discussion

We found no evidence supporting the hypothesis that predation of lionfish on small prey individuals (Albins and Hixon 2008, Green et al. 2012, Albins 2013) affects adult reef fish communities. Across sixteen fore-reef sites on the MBR in Belize, there was no relationship between the abundance of lionfish and the abundance, species richness, diversity, or community composition of all native fishes or native prey species. The only factors significantly related to fish community structure were reef complexity, site protection status, and time since the lionfish invasion. Habitat complexity not only influences reef fish communities through habitat availability (Friedlander et al. 2003), but can also reduce mortality due to predation (Beukers and Jones 1998). On the MBR in Belize, habitat complexity was more important in structuring reef fish communities than predation by lionfish, but did not modulate the effect of lionfish (i.e. the interaction between lionfish abundance and habitat complexity had no significant effect on reef fish community

structure). Total fish diversity was lower on protected sites compared to unprotected, which is the opposite of the expected effect of fishing regulations on reef fish communities (Friedlander et al. 2003). This difference in diversity is likely indicative of differences in species evenness, however, as species richness was not influenced by protection status. For example, differences in fishing regulations may influence the relative abundances of reef fish species, resulting in communities where species evenness, and therefore diversity, is lower on protected sites compared to unprotected sites. Although there was a significant effect of time since invasion on total fish diversity, this pattern was not related to lionfish abundance. Additionally, reef fish community composition changed significantly on some sites, but there was no overall shift in composition from pre- to post-invasion across the MBR. Lionfish abundance had no significant effect on reef fish community structure or composition, so these changes over time are not due to the lionfish invasion. Instead, this is likely indicative of inter-annual fluctuations in the reef fish community responses, potentially due to stochastic population variation, recruitment success, or variability in unmeasured environmental drivers.

The specific effects of an invasive predator on native communities (i.e., declines in abundance/diversity or changes in community composition) can be estimated with controlled lab or field experiments. However, to properly inform management and conservation strategies, it is necessary to determine the importance of the invasive predator relative to existing forces structuring native communities including other anthropogenic impacts. Prey populations are naturally regulated by bottom-up forces such as habitat availability, top-down forces such as predation by native species, inter- and intra-specific competition, as well as stochastic processes such as variability in recruitment or environmental suitability (Hunter

and Price 1992, Power 1992). Additionally, anthropogenic stressors including exploitation, habitat degradation and fragmentation, pollution, and climate change are threatening natural communities (Halpern et al. 2008). These natural and anthropogenic forces could be more important for community structure than the effect of the invasive predator, suggesting conservation efforts should be focused on threats other than the invader. However, it is difficult to design controlled experiments that include the suite of factors influencing prey communities. Additionally, the relative importance of recruitment vs. post-recruitment processes cannot necessarily be extrapolated from isolated habitat patches to contiguous habitats in the marine environment (Ault and Johnson 1998). Our study design allowed us to use the lionfish invasion on the MBR as an “accidental” large-scale experiment to quantify the effects of invasive lionfish over time within a natural reef system. Our goal was to determine if the effect of lionfish was apparent in reef fish communities within the context of the existing environmental and annual variability across our fore-reef sites. We found no evidence to support this. Instead, reef fish communities were structured by bottom-up effects of habitat complexity and stochastic processes (e.g., annual variability) rather than the top-down pressure of predation by lionfish on reef fish recruits.

Scale-dependent mechanisms may also be relatively more important in structuring reef fish communities compared to the predation of lionfish. The existing documentation of negative effects of lionfish on prey individuals comes from experimental reefs near Lee Stocking Island, Bahamas, covering an area of $\sim 6 \text{ km}^2$ and a total surveyed area of $\sim 80 \text{ m}^2$ (20 $\sim 4 \text{ m}^2$ reefs) (Albins and Hixon 2008, Albins 2013) as well as one observational study on a single continuous reef with sites separated by 1 km in New Providence, Bahamas (Green et al. 2012). In contrast, our study tested the effect of the lionfish invasion across a latitudinal

scale, covering a ~250 km stretch of reef and a total surveyed area of ~48 km² (~6 transects of 50 x 10 m area on 16 sites) including 16 different sites along the MBR in Belize Fig. 1, Table S1). In our large-scale and complex study system, reefs may be connected (Tang et al. 2006, Soto et al. 2009) and influenced by meta-population dynamics (Kritzer and Sale 2004). The underlying impact of lionfish on individual reefs could be counteracted by recruitment or immigration from surrounding reefs within the system (i.e., the “rescue effect” (Gotelli 1991)) depending on reef connectivity (Ault and Johnson 1998). Potential evidence of this is provided in the lionfish removal experiment by Green et al. (2014) where the effect of lionfish on small prey biomass differed between low and high lionfish density treatments, yet this pattern was not apparent in the biomass of larger individuals between treatments. Instead, the biomass of larger individuals increased across all lionfish densities treatments, which authors attributed to inter-reef movement of individuals offsetting the underlying effect of lionfish (Green et al. 2014). Isolated study systems, such as those used in previous studies (Albins and Hixon 2008, Albins 2013), are essential for experimentally quantifying the effect of an *in-situ* treatment (e.g., the presence of lionfish). However conclusions from such experiments may not necessarily be applicable to more complex systems without further investigation. Our study design allowed us to determine that the effect of lionfish on reef fish communities was not apparent within a complex reef system.

The effects of invasive species may be influenced by the composition or structure of the recipient communities, especially in terms of potential competition pressure (i.e., the “biotic resistance hypothesis”(Elton 1958)). One possible explanation for our results may be that competitor abundance declined as lionfish abundance increased during the time of our study, so that net top-down predation pressure on recruits remained fairly constant (although

lionfish have been shown to have a greater effect on fish recruits than native predators on experimental reefs (Albins 2013)). However, this is not the case on our sites as the abundance of piscivores that are ecologically similar to lionfish (species listed in Table S4) did not decrease across our sites from 2009 to 2013 (Fig. S4). Instead, average competitor abundance remained fairly constant, with a slightly positive trend over time (Fig. S4). Therefore, the lack of apparent effect of lionfish on reef fish communities is likely not due to changes in native piscivore abundance. However, the effect of lionfish may be limited by the relative abundance of native predators. Native predators do not control lionfish (Hackerott et al. 2013, Valdivia et al. 2014), but proportionally lower lionfish biomass in comparison to native predators may limit the relative influence of lionfish on reef fish communities. On impacted reefs, lionfish can comprise a sizable portion of the total fish predator biomass, becoming one of the most important predators. For example, lionfish were reported to increase from 23% to 40% of the total predator biomass on reefs in New Providence, Bahamas from 2008-2010, coinciding with a decrease in biomass of small prey individuals (Green et al. 2012). During experiments that demonstrated a significant effect on reef fish recruits, lionfish comprised ~50-100% of the total predator biomass (Albins and Hixon 2008, Albins 2013). In contrast, lionfish biomass in our study peaked at ~27% of the total biomass of ecologically similar predators (Table S4) in 2011 and remained relatively low, ~10-11%, during 2012 and 2013 (Fig. S5). While previous studies showed that lionfish can control reef fish recruits when acting as one of the most important, or the only, predators, it is likely that the effect of lionfish on reef fish communities is context-dependent.

The effect of lionfish on reef fish communities may also be density-dependent. On patch reefs, the effect of lionfish on the biomass of small prey individuals can be limited if

lionfish densities are reduced below a reef-specific threshold (Green et al. 2014). The lionfish densities across our sites post-invasion (2012 and 2013) were significantly lower (one-tailed t-test; $t = 8.3$; $p\text{-value} = 2.4e^{-6}$) than the lionfish densities (mean \pm SE; $\sim 138.9 \pm 14.7$ individuals/ha) in the 25th percentile of the thresholds predicted for patch reefs in Eleuthera, Bahamas (Green et al. 2014). While the values of threshold densities vary across different communities, site-specific “thresholds” for our larger fore-reef sites are more likely to be higher rather than lower than those estimated for smaller patch reefs, as reef fish communities may be more strongly structured by recruitment on isolated patch reefs than on contiguous reef systems (Ault and Johnson 1998). Therefore, it is likely that the lionfish densities on our sites were lower than the site-specific “threshold” levels, which could explain why the predicted effects of lionfish were not apparent in our study.

The density of lionfish also differed between our study and previous studies testing the effects of lionfish on small prey individuals. On experimental reefs in Lee Stocking Island, Bahamas, lionfish treatments included one lionfish per $\sim 1 \text{ m}^2$, 3 m^2 , and 4 m^2 reefs which scales to unrealistic densities of $\sim 2500\text{-}10,000$ individuals/ha (Albins and Hixon 2008, Albins 2013) which is 1-2 orders of magnitude higher than the maximum recorded density (Green and Côté 2009) and $\sim 100\text{-}900$ times higher than our average post-invasion densities. Again, such controlled experiments are essential to determine the *potential* effect of a treatment (e.g., the presence of lionfish). However, it is important to consider the applicability of the experimental treatment to existing systems and to measure the *realized* effects at realistic treatment levels. In New Providence, Bahamas, the existing levels of lionfish abundance were reported to coincide with a decrease in prey biomass, although these patterns were not analyzed together at the transect or site level (Green et al. 2012). While the

values of lionfish densities were not reported in that study, values reported on sites in New Providence during the time of the study (101.6 ± 28.6 individuals/ha in 2008 and 87.9 ± 24.2 individuals/ha in 2010 (Darling et al. 2011, Valdivia et al. 2014)) were significantly higher (one-tailed t-test; $t=4.2$; $p\text{-value}=2.0e-4$) than lionfish densities across our sites post-invasion (2012 and 2013). Additionally, maximum lionfish densities reached 393.3 ± 83.4 individuals/ha in 2008 on three sites in New Providence, Bahamas (Green and Côté 2009) while the highest lionfish density on our study sites was 70 ± 29.2 individuals/ha at Pompion in 2012. Even 4 years after the first lionfish was reported in each region (New Providence: 2005 and Belize: 2009 (Schofield 2009)), lionfish densities were significantly higher (one-tailed t-test; $t=3.1$; $p\text{-value}=4.0e-3$) in New Providence, Bahamas than on our sites in Belize, so time since invasion does not explain the difference in densities. It is possible that lionfish propagule pressure is lower in Belize compared to the Bahamas due to higher spatial separation and/or lower connectivity with the point of origin of the invasion (Southern Florida (Schofield 2009)), similar to patterns documented in terrestrial invasions (Hastings et al. 2005, Alston and Richardson 2006).

Lionfish densities in Belize are likely controlled by a combination of natural (e.g., propagule pressure, environmental suitability, etc.) and anthropogenic forces (e.g., lionfish removals by reef managers and recreational divers), although the relative influence of these factors on lionfish in Belize is unknown. While an effect of lionfish may become apparent in the future (due to increases in lionfish density or decreases in prey productivity), current densities of lionfish have had no apparent effect on reef fish communities on our sites. It seems that the current level of control on lionfish densities has been sufficient to prevent impacts, therefore *additional* lionfish removal efforts may not be necessary on our sites.

Additionally, we suggest quantifying the importance of current removal efforts, relative to natural forces, on both lionfish densities and impacts to determine the necessary level of lionfish removals. This can increase reef management efficiency, which is essential in countries such as Belize where reef managers are often resource-limited. Lionfish densities and apparent impacts should also be monitored on our sites through time, as well as across other reef sites and different habitat types in Belize, to identify sites where targeted removal efforts may be necessary. It is unlikely that lionfish can be completely extirpated from invaded reefs, but our study suggests this may not be necessary to prevent impacts on reef fish communities. Therefore, future efforts should be focused on monitoring for apparent impacts of lionfish so that management resources are not wasted on extraneous lionfish removals in areas where they may not be necessary.

Our study provides the first case study quantifying the effects of lionfish on reef fish communities that may be applicable across the Caribbean. Lionfish densities on our sites in Belize are more comparable to published densities on reefs in Mexico, Cuba, The Bahamas, Colombia, and Venezuela than densities reported in New Providence, Bahamas (Fig. 5, Text S1), previously assumed to be representative of the entire Caribbean. The effects of lionfish are likely density dependent (Green et al. 2014) so the effect of lionfish in Belize may be representative of the effect in areas with similar densities. Additionally, lionfish densities in these areas across the Caribbean were below the estimated “effect threshold” and comparable to densities reported in the native range (Fig. 5, Text S1) which may further support the possibility that current lionfish densities in these areas might not be detectably changing reef fish communities. However, effects of lionfish are also dependent upon prey productivity so

lionfish densities and apparent impacts should be continually monitored on these coral reef sites as well as on additional sites and habitats across the Caribbean.

Conclusions

Caribbean coral reefs face a growing list of current and future threats including overfishing of predatory fish (Stallings 2009), exploitation of herbivores (Mumby 2006), other local anthropogenic stressors (Mora 2008), as well as the stresses associated with global climate change (Hoegh-Guldberg et al. 2007). For reef managers tasked with conserving these ecologically and economically important ecosystems, conservation priorities must be applicable within the context of existing variability and stresses across reefs. Our study is the first to quantify the effects of lionfish on reef fish communities across a large-scale, complex reef system, within the context of existing variability. While lionfish were predicted to decimate native reef fish communities (Green et al. 2012, Albins and Hixon 2013) and have the potential to influence small prey individuals in experimental settings (Albins and Hixon 2008, Albins 2013), lionfish have had no apparent effect on reef fish communities in Belize. The level at which lionfish are currently controlled in Belize, due to lionfish culling or other factors, have been sufficient to prevent detectable changes in reef fish communities. Therefore reef managers in Belize may not need to increase their efforts to control lionfish. This management recommendation may be applicable to other regions in the Caribbean with comparable lionfish densities. Lionfish removal efforts are common throughout the Caribbean and can successfully reduce the abundance (Barbour et al. 2011, Frazer et al. 2012) and effects (Green et al. 2014) of lionfish. However, our study suggests that the management effort necessary to prevent the impacts of lionfish may not be as generalizable as previously assumed, but instead will likely vary across the Caribbean and

depend upon local conditions and lionfish densities. The densities and effects of lionfish should be monitored to identify sites where targeted removals may be necessary to conserve native reef fish communities as well as to identify sites that should be more strenuously managed for threats other than lionfish. While the lionfish invasion is a Caribbean-wide problem, the development of management plans and priorities should be region-specific.

FIGURES

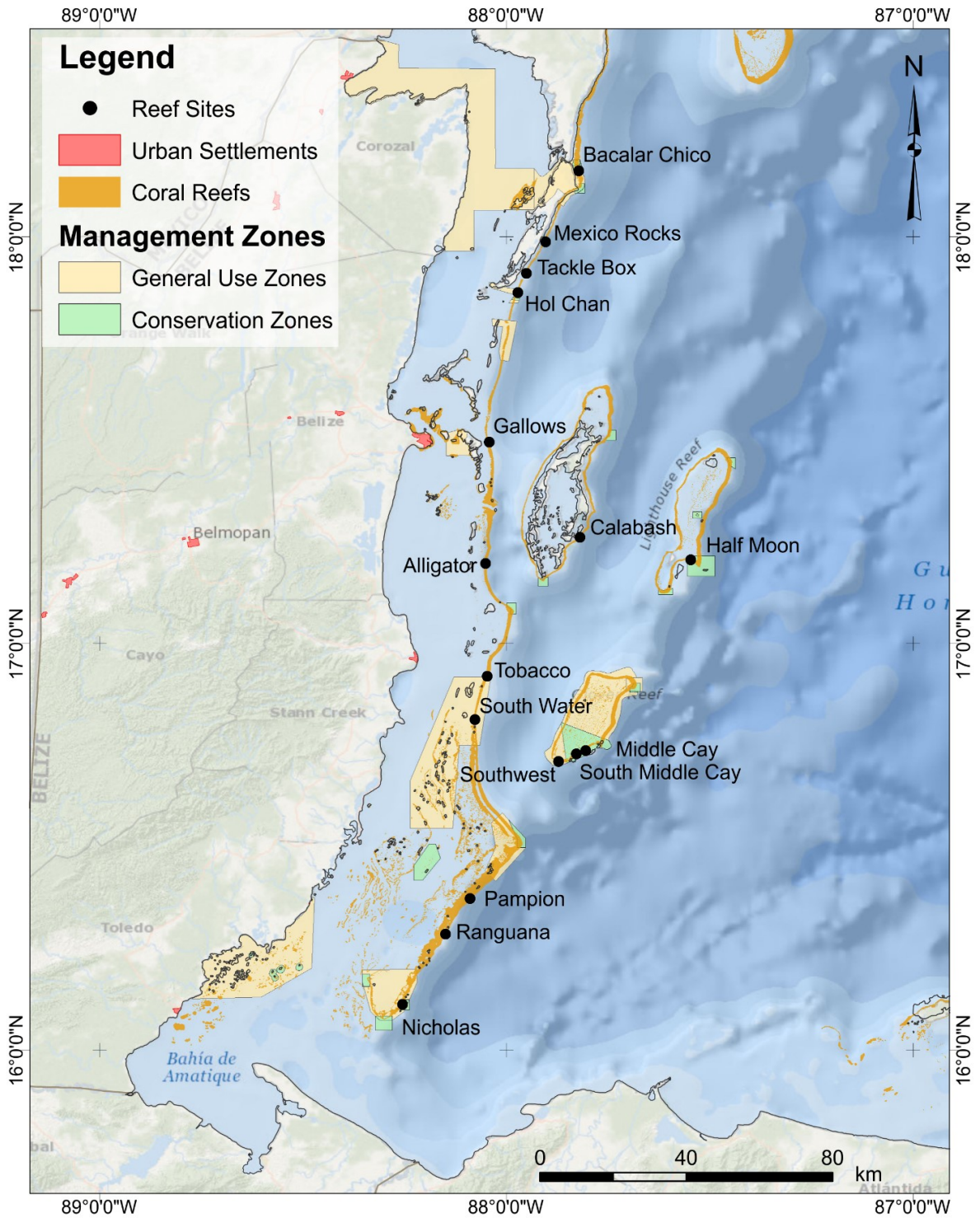


Figure 1. Location of Survey Sites. Location of surveys sites. For sites abbreviations, coordinates, and other site info, refer to Table S1.

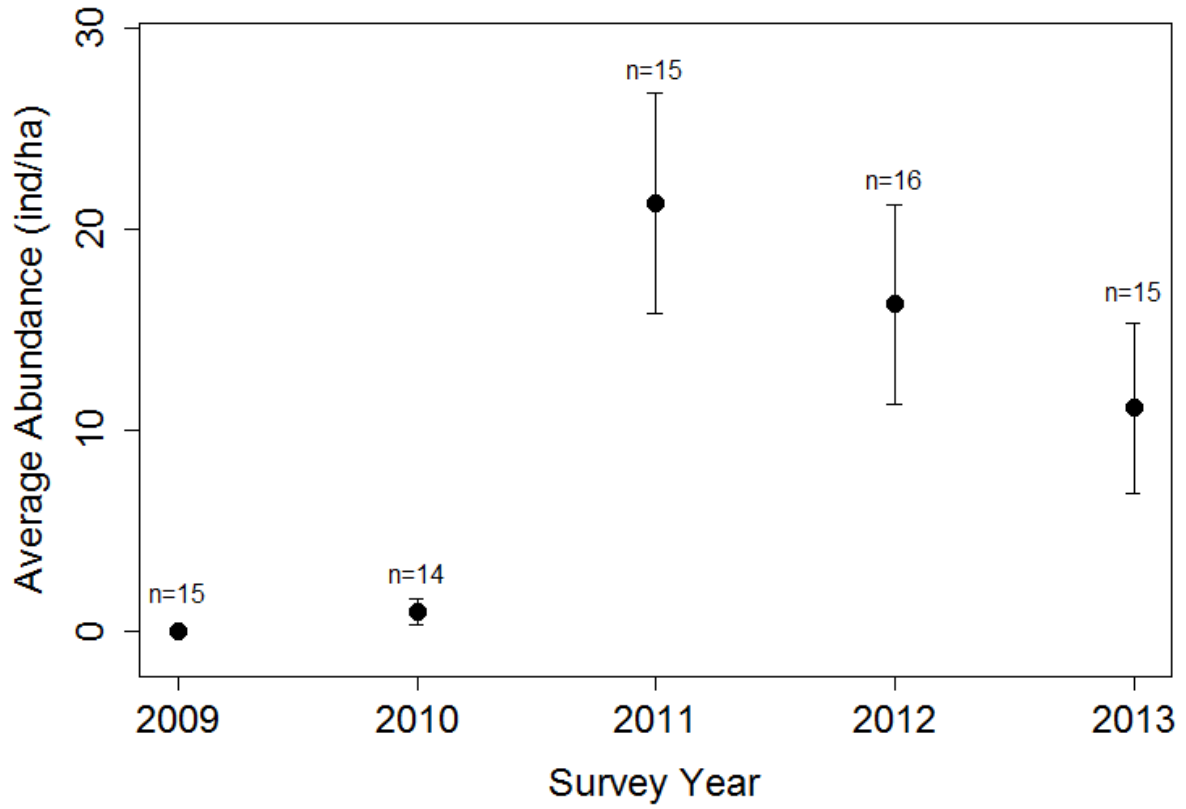


Figure 2. Average Lionfish Abundance from 2009 to 2013. Lionfish abundance (individuals/ha), averaged over all survey sites for each survey year. N is the number of sites surveyed in each year.

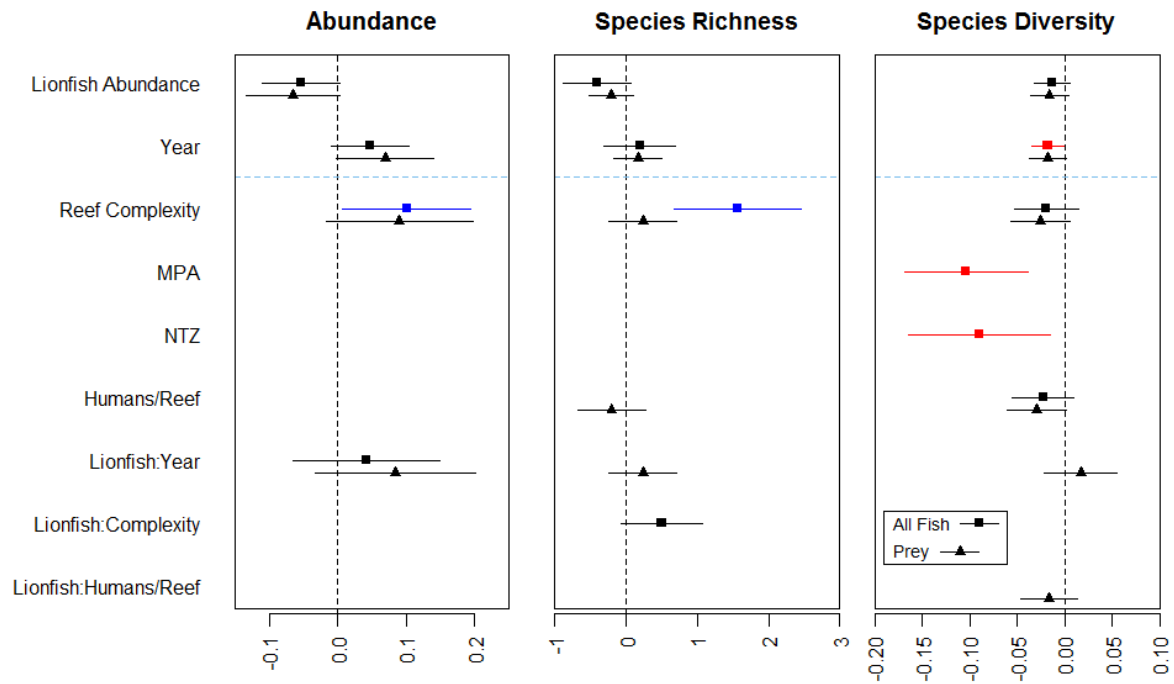


Figure 3. Coefficient Estimates (mean \pm 95% confidence interval) **for Each Model.** Total fish and prey fish abundance, species richness, species diversity were each modeled with the predictors of interest (lionfish abundance and years since the lionfish invasion) as well as the abiotic covariates (reef complexity, protection status, humans/reef) and possible interactions, separated by the dotted blue line. The coefficient estimates for the models of total fish are indicated by squares while those of prey fish are indicated by triangles. Significantly positive estimates are shown in blue and significantly negative in red (details in Table S3).

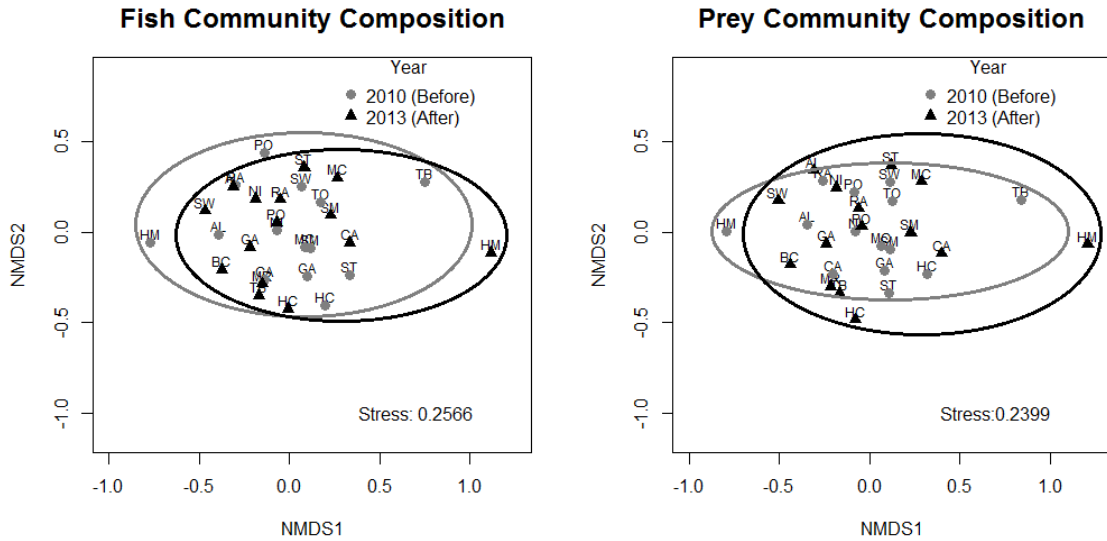


Figure 4. Total Fish and Prey Fish Community Composition in 2010 and 2013. Community composition of total fish and prey fish communities within two-dimensional space using nonmetric multidimensional scaling (NMDS) of Bray-Curtis dissimilarity at the site level. Communities in 2010 (before invasion) are shown in gray dots and circled in grey while communities in 2013 (after invasion) are shown in black triangles and circled in black.

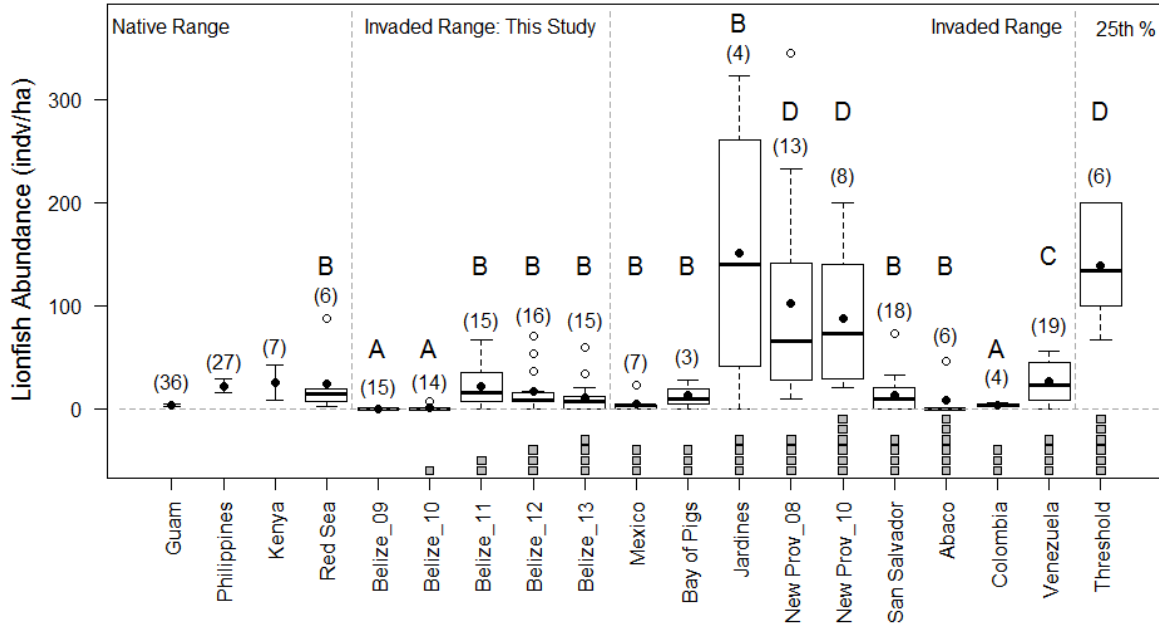


Figure 5. Lionfish Abundance in Native and Invaded Ranges and Estimated Impact Threshold. Lionfish abundance (individuals/ha) in the native range (left panel) and the invaded range (center two panels), compared with the 25th percentile abundance threshold predicted to limit effects of lionfish on reef fish (Green et al. 2014) (right panel); details in Text S1. Boxplots were used to represent lionfish abundance if site-level data was available. If not, averages and confidence intervals were plotted. The number of sites surveyed in each area are in parentheses. Average lionfish abundance for each area is represented by a solid black dot. Letters indicate significant differences ($p < 0.05$) in lionfish abundance across all studies where site-level data was available. The number of years since the region-specific onset of invasion (Schofield 2009) at the time of each study are represented by the number of gray boxes along the bottom panel.

APPENDIX 1: SUPPLEMENTAL TABLES

Table S1. Site Information and Coordinates.

Site Name	Site Code	Latitude (N)	Longitude (W)	Prot. Level	Prot. Status	Reef Complexity	Humans/ Reef
Alligator	AL	17.1966	-88.0512	None	No	2	6.39
Bacalar Chico	BC	18.16282	-87.8222	NTZ	Yes	2.92	4.53
Calabash	CA	17.26147	-87.8197	None	No	2.42	14.55
Gallows	GA	17.49592	-88.0426	None	No	3.5	5.9
Hol Chan	HC	17.86343	-87.9724	NTZ	Yes	2	14.03
Half Moon	HM	17.2056	-87.5468	NTZ	Yes	5	3.28
Middle Cay	MC	16.73703	-87.8054	MPA	Yes	1.67	0.73
Mexico Rocks	MR	17.98782	-87.9038	None	No	2.58	11.63
Nicholas	NI	16.1123	-88.2559	MPA	Yes	2.33	11.81
Pampion	PO	16.3731	-88.0891	None	No	2.92	2.61
Ranguana	RA	16.28501	-88.1503	None	No	2.83	2.79
South Middle Cay	SM	16.72875	-87.8287	MPA	Yes	2	1
Southwest	ST	16.11247	-88.2711	MPA	Yes	2.33	7.8
South Water	SW	16.81346	-88.0776	MPA	Yes	2.08	2.45
Tackle Box	TB	17.91056	-87.9508	None	No	2.75	12.86
Tobacco	TO	16.91911	-88.0476	None	No	2.08	4.01

Table S2. Prey Species and Sources. Fish species that are documented prey for lionfish in the Caribbean.

Family	Species Name	Sources
Acanthuridae	<i>Acanthurus bahianus</i>	3
Acanthuridae	<i>Acanthurus chirurgus</i>	6,8
Acanthuridae	<i>Acanthurus coeruleus</i>	5
Aulostomidae	<i>Aulostomus maculatus</i>	2,3,4,7
Cirrhitidae	<i>Amblycirrhitus pinos</i>	3
Gobiidae	<i>Coryphopterus dicrus</i>	3,5,7
Grammatidae	<i>Gramma loreto</i>	1,2,3,4,7
Haemulidae	<i>Haemulon aurolineatum</i>	7
Haemulidae	<i>Haemulon flavolineatum</i>	1,7
Holocentridae	<i>Holocentrus rufus</i>	2,7
Labridae	<i>Bodianus rufus</i>	3,4,6,7,8
Labridae	<i>Clepticus parrae</i>	2,3,4,7
Labridae	<i>Halichoeres bivittatus</i>	2,3,4,7,8
Labridae	<i>Halichoeres garnoti</i>	1,2,3,4,5,7,8
Labridae	<i>Halichoeres maculipinna</i>	3,4,5,6
Labridae	<i>Halichoeres pictus</i>	3,5,6
Labridae	<i>Thalassoma bifasciatum</i>	1,2,3,4,6,7,8
Mullidae	<i>Pseudupeneus maculatus</i>	3,4,7
Pomacentridae	<i>Abudefduf saxatilis</i>	1,8
Pomacentridae	<i>Chromis cyanea</i>	2,3,4,7

Pomacentridae	<i>Chromis insolata</i>	3
Pomacentridae	<i>Chromis multilineata</i>	2,3,4,7
Pomacanthidae	<i>Holacanthus ciliaris</i>	6
Pomacentridae	<i>Stegastes leucostictus</i>	3,6,8
Pomacentridae	<i>Stegastes partitus</i>	1,2,3,4,5,6,7
Pomacentridae	<i>Stegastes variabilis</i>	2,3,4,5,7
Scaridae	<i>Scarus iserti</i>	1,3,7
Scaridae	<i>Scarus taeniopterus</i>	1
Scaridae	<i>Sparisoma atomarium</i>	5,6
Scaridae	<i>Sparisoma aurofrenatum</i>	1,2,4,5,6,7
Scaridae	<i>Sparisoma viride</i>	1,3,5,6
Scorpaenidae	<i>Scorpaena plumieri</i>	6
Serranidae	<i>Cephalopholis cruentata</i>	1,2,4,7
Serranidae	<i>Epinephelus striatus</i>	3,4
Serranidae	<i>Serranus tigrinus</i>	2,3,4,5,6,7
Tetraodontidae	<i>Canthigaster rostrata</i>	3,5,6,7

Sources: 1. Valdez-Moreno et al (2012), 2. Cote et al (2013), 3. Morris and Akins (2009), 4. Green et al (2012), 5. Albins and Hixon (2008), 6. Albins (2013), 7. Green and Cote (2014), 8. Layman and Allgeier (2011)

Table S3. Coefficient Estimates and p-values for Each Model. Coefficient estimates and p-values for each model of (A) total fish and (B) prey fish abundance, species richness, and species diversity. Significant coefficient estimates are shown in bold at alpha 0.05. “NA” indicates that the predictor did not remain in the final model following model averaging.

Table S3A. All Fish	Total Abundance		Species Richness		Species Diversity	
	Estimate	p-value	Estimate	p-value	Estimate	p-value
Lionfish Abundance	-0.0530	0.0632	-0.4051	0.0935	-0.0130	0.1581
Year	0.0481	0.0931	0.1904	0.4460	-0.0177	0.0411
MPA	NA	NA	NA	NA	-0.1036	0.0015
NTZ	NA	NA	NA	NA	-0.0897	0.0177
Reef Complexity	0.1009	0.0328	1.5639	0.0005	-0.0192	0.2522
Humans/Reef	NA	NA	NA	NA	-0.0227	0.1621
Lionfish Abundance : Year	0.0422	0.4367	NA	NA	NA	NA
Lionfish Abundance : Protected	NA	NA	NA	NA	NA	NA
Lionfish Abundance : Reef Complexity	NA	NA	0.5012	0.0773	NA	NA
Lionfish Abundance : Humans/Reef	NA	NA	NA	NA	NA	NA

Table S3B. Prey Fish	Total Abundance		Species Richness		Species Diversity	
	Estimate	p-value	Estimate	p-value	Estimate	p-value
Lionfish Abundance	-0.0656	0.0555	-0.2097	0.177	-0.0159	0.1183
Year	0.0695	0.0514	0.1656	0.324	-0.0180	0.0649
MPA	NA	NA	NA	NA	NA	NA
NTZ	NA	NA	NA	NA	NA	NA
Reef Complexity	0.0900	0.0953	0.2372	0.324	-0.0253	0.1041
Humans/Reef	NA	NA	-0.2016	0.392	-0.0294	0.0619
Lionfish Abundance : Year	0.0848	0.1507	0.3473	0.241	0.0170	0.3787
Lionfish Abundance : Protected	NA	NA	NA	NA	NA	NA
Lionfish Abundance : Reef Complexity	NA	NA	NA	NA	NA	NA
Lionfish Abundance : Humans/Reef	NA	NA	NA	NA	-0.0163	0.2696

Table S4. Competitor Species. Fish species present on our surveys that are ecologically similar to lionfish based on diet from Green et al. (2012) and FishBase.

Family	Species Name
Aulostomidae	<i>Aulostomus maculatus</i>
Carangidae	<i>Alectis ciliaris</i>
Carangidae	<i>Carangoides ruber</i>
Carangidae	<i>Caranx bartholomaei</i>
Carangidae	<i>Caranx crysos</i>
Carangidae	<i>Elagatis bipinnulata</i>
Lutjanidae	<i>Lutjanus analis</i>
Lutjanidae	<i>Lutjanus apodus</i>
Lutjanidae	<i>Lutjanus cyanopterus</i>
Lutjanidae	<i>Lutjanus jocu</i>
Lutjanidae	<i>Lutjanus mahogoni</i>
Lutjanidae	<i>Lutjanus synagris</i>
Scorpaenidae	<i>Scorpaena plumieri</i>
Serranidae	<i>Cephalopholis cruentata</i>
Serranidae	<i>Cephalopholis fulva</i>
Serranidae	<i>Epinephelus adscensionis</i>
Serranidae	<i>Epinephelus guttatus</i>
Serranidae	<i>Epinephelus itajara</i>
Serranidae	<i>Epinephelus morio</i>
Serranidae	<i>Epinephelus striatus</i>

Serranidae	<i>Mycteroperca bonaci</i>
Serranidae	<i>Mycteroperca interstitialis</i>
Serranidae	<i>Mycteroperca tigris</i>
Serranidae	<i>Mycteroperca venenosa</i>

APPENDIX 2: SUPPLEMENTAL FIGURES

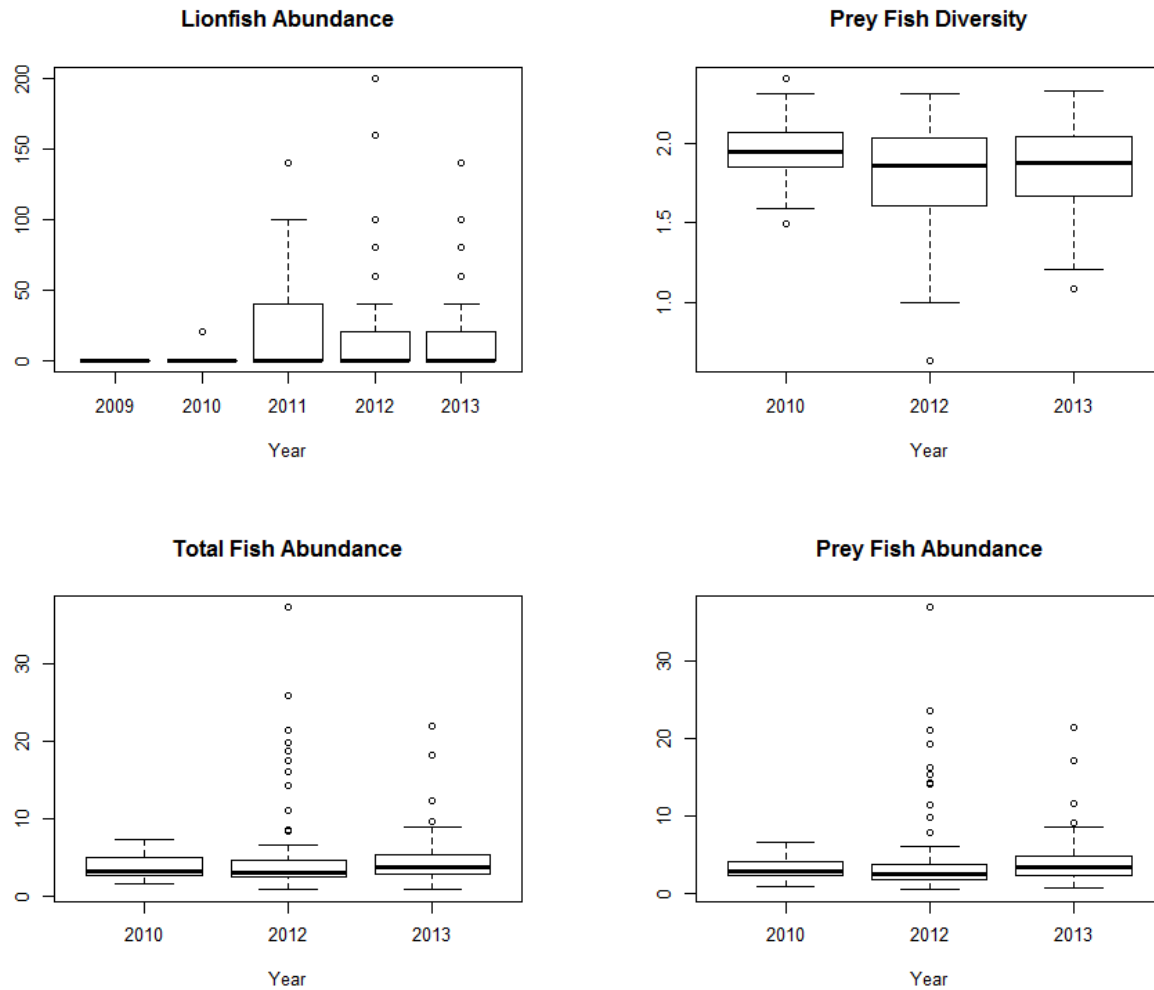


Figure S1. Boxplots of Lionfish Abundance, Prey Fish Diversity, and Total and Prey Fish Abundance. Boxplots were used to determine the outlier transects that were removed from lionfish abundance (individuals/ha) in all models and from prey fish diversity and total and prey fish abundance (individuals/m²) in each respective model.

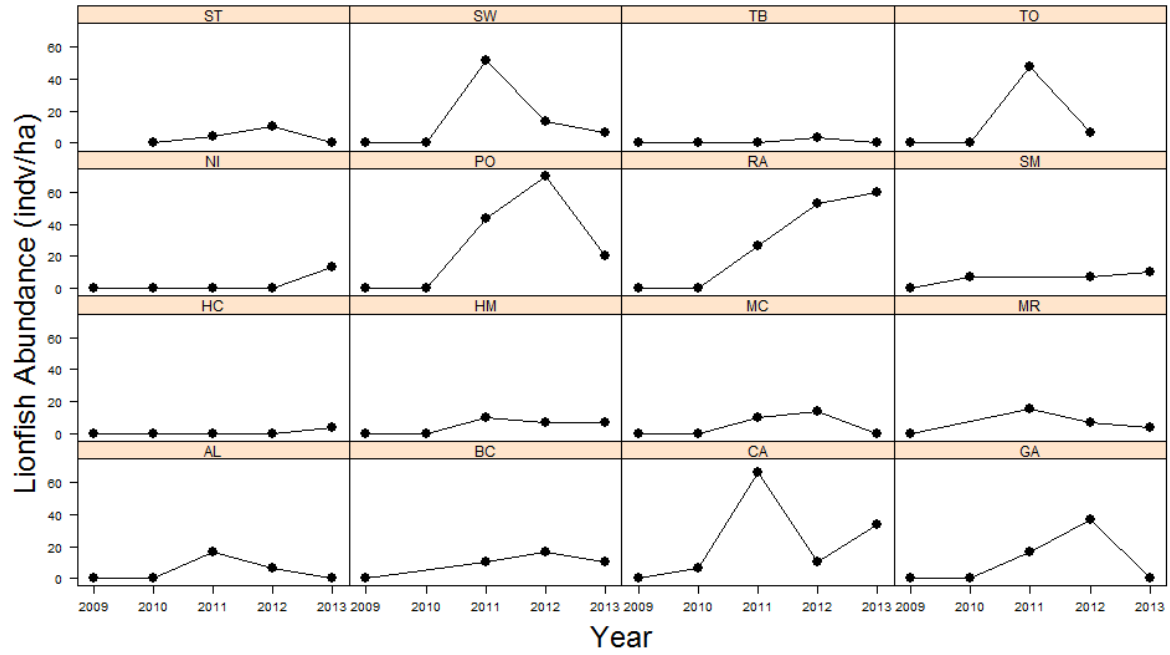


Figure S2. Lionfish Abundance from 2009 to 2013 on Each Site. Lionfish abundance (individuals/ha) over survey years on each survey site.

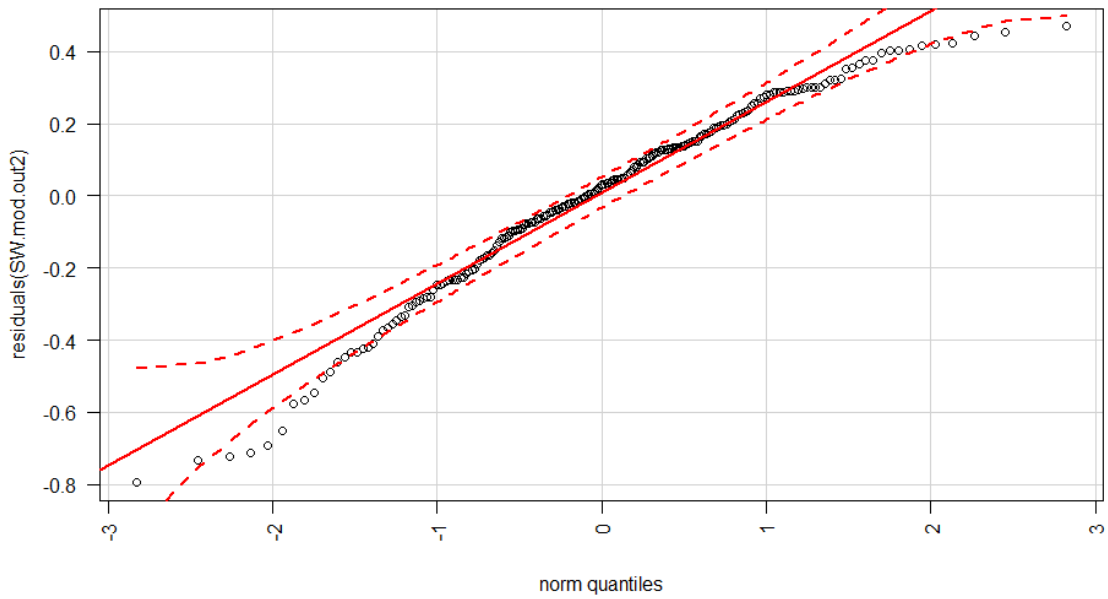


Figure S3. Quantile-Quantile Plot of the Residuals of the Model of Total Fish Species Diversity. Residuals of the models of total fish species diversity are slightly non-normal due to a few values at the extremes.

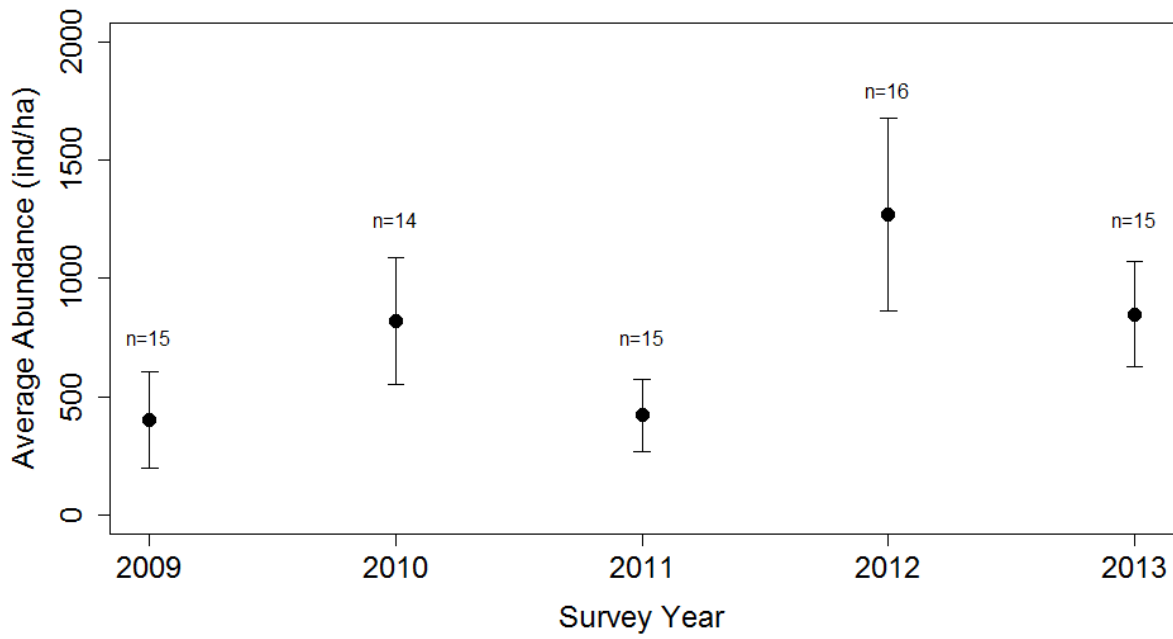


Figure S4. Abundance of Native Competitors 2009-2013. The abundance (individuals/ha) of native piscivores that are ecologically similar to lionfish (see Table S4) from 2009-2013, averages across site. N is the number of sites surveyed in each year.

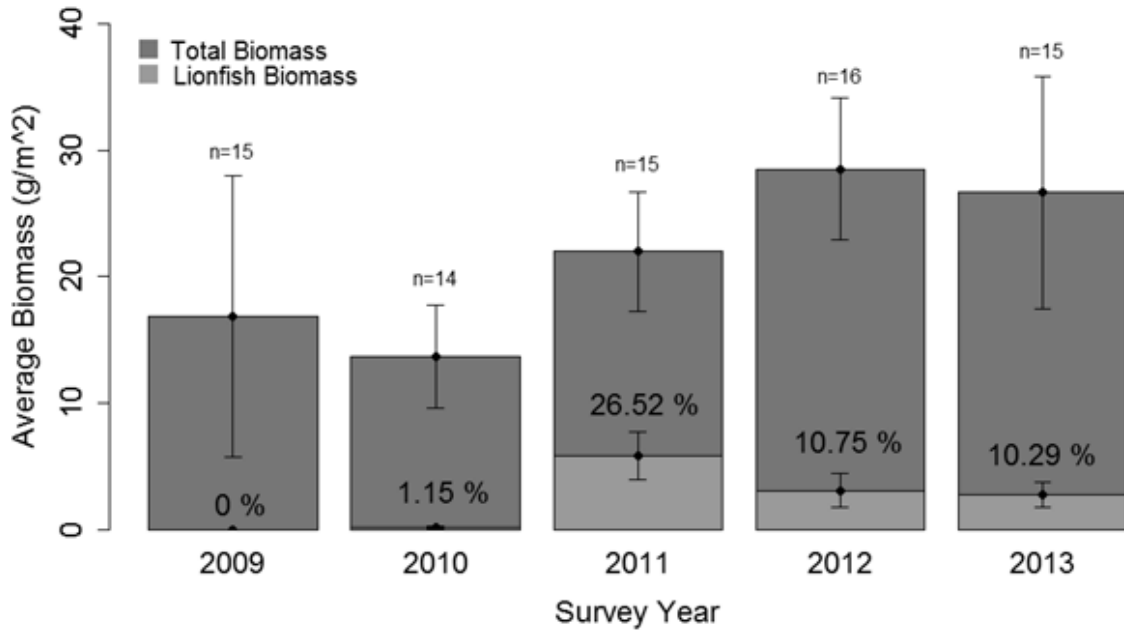


Figure S5. Biomass of Lionfish and Native Competitors from 2009 to 2013. The biomass (g/m^2) of lionfish is shown in light gray, and the total biomass of all ecologically similar native piscivores (see Table S4) is shown in dark gray, from 2009-2013, averaged across sites. The percentage of total predator biomass contributed by lionfish is shown above the average biomass of lionfish in each year. N is the number of sites surveyed in each year.

APPENDIX 3: SUPPLEMENTAL TEXT

Text S1. Published Values of Lionfish Densities in Native and Invaded Range.

Lionfish density data was collected from published studies with methods comparable to ours. Only studies surveying coral reef habitats with belt transects of defined lengths and widths were included. Additionally, only studies that presented averages \pm standard deviation/error, or site-level data were included. Lionfish densities from a total of eight published studies, representing two regions of the Indo-Pacific (native range) and five regions of the Caribbean (invaded range), were compared to our values from Belize. Lionfish densities in Belize were not significantly different than those measured in Mexico (two-tailed t-test; $t=-2.0$; $p\text{-value}=6.4e^{-2}$), Bay of Pigs, Cuba (two-tailed t-test; $t=-0.1$; $p\text{-value}=0.9$), Jardines de la Riena, Cuba (two-tailed t-test; $t=-1.9$; $p\text{-value}=0.1$), Abaco, Bahamas (two-tailed t-test; $t=-0.7$; $p\text{-value}=0.5$) (Valdivia et al. 2014), and San Salvador, Bahamas (two-tailed t-test; $t=-0.2$; $p\text{-value}=0.9$) (Anton et al. 2014), and were significantly higher than those in Colombia (one-tailed t-test; $t=-3.0$; $p\text{-value}=2.3e^{-3}$) (Bayraktarov et al. 2014) (Fig. 5). Lionfish densities in Venezuela (Agudo and Salas 2014) were higher than those in Belize (one-tailed t-test; $t=-2.2$; $p\text{-value}=1.8e^{-3}$), but significantly lower than those reported in New Providence, Bahamas (one-tailed t-test; $t=3.5$; $p\text{-value}=1.9e^{-3}$) (Darling et al. 2011, Valdivia et al. 2014) (Fig. 5). Furthermore, lionfish densities on our sites in Belize were not significantly different (two-tailed t-test; $t=0.8$; $p\text{-value}=0.5$) in the native range, specifically the Red Sea where site-level data was available (McTee and Grubich 2014), and were comparable to densities in Guam and the Philippines (Cure et al. 2014), as well as Kenya (Darling et al. 2011) (Fig. 5).

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