

1 **Fishing degrades size structure of coral reef fish communities**

2 **SUPPORTING INFORMATION**

3 **Appendix S1**

4 **Explanatory covariate processing**

5 *Human covariates*

6 No standard measure of fishing effort was available across all islands sampled,
7 so instead we quantified exploitation pressure using two metrics of broad human
8 impacts on coral reef fish communities. Higher human population densities (Graham
9 et al., 2005; Williams et al., 2011; Cinner et al., 2012; MacNeil et al., 2015; Williams
10 et al., 2015) and greater access to markets (Brewer et al., 2012; Cinner et al., 2012)
11 have been associated with degraded reef fish assemblages, primarily due to increased
12 fishing effort. We extracted 2010 estimates of population density for a 20 km buffer
13 (SEDAC) and divided this by the forereef area (km²). Following Cinner et al. (2012),
14 we estimated the shortest distance (km) from each site to the provincial capital as a
15 proxy of market access (using ArcGIS).

16

17 *Environmental covariates*

18 We extracted the average minimum SST (°C) from the National
19 Oceanographic Data Center's Coral Reef Temperature Anomaly Database (CoRTAD)
20 based on AVHRR Pathfinder data on a weekly time-scale between 1982-2008 at a
21 ~4.6 x 4.6 km resolution (<http://www.nodc.noaa.gov/SatelliteData/Cortad>). Net
22 primary productivity (mg C m⁻² day⁻¹) was modelled by NOAA CoastWatch based on
23 satellite measurements of photosynthetically available radiation (NASA's SeaWiFS),
24 SST (NOAA's National Climatic Data Center Reynolds Optimally-Interpolated SST),
25 and chlorophyll *a* concentration (NASA Aqua MODIS) estimated every 8 days

26 between 2002-2013 at a ~4.6 x 4.6 km resolution
27 (<http://coastwatch.pfeg.noaa.gov/erddap/griddap/erdPPbfp28day.graph>). To avoid
28 introducing bias from increased reflectance in shallow waters, we used bathymetry
29 data from the STRM30_plus high-resolution (~1 x 1 km) bathymetry dataset
30 (http://topex.ucsd.edu/WWW_html/srtm30_plus.html) and a focal productivity cell
31 was excluded if any bathymetry cell within its bounds was <30 m depth (Gove et al.,
32 2013). We then estimated the mean SST value and mean productivity value per site
33 by averaging SST or productivity values across the cell that included the site (if not
34 excluded based on depth) and the closest 3 neighbouring cells within 9.3 km of either
35 the shoreline or site that also passed the depth filter.

36 The extent of coral reef habitat around each site was obtained by merging data
37 from two primary sources, the Millennium Coral Reef Mapping project (data accessed
38 from the University of South Florida Institute for Marine Remote Sensing and
39 through the UNEP World Conservation Monitoring Centre, Andréfouët et al., 2005)
40 and coral maps for Asian Pacific region produced under the “Coral Reef Habitat
41 Map” project by the Japanese Ministry of the Environment
42 (http://coralmap.coremoc.go.jp/sangomap_eng/; accessed 4/28/2011). We merged all
43 non-land geomorphological types mapped in the Millennium data as this classification
44 best matched the known extent of reef habitat based on the fish survey locations.
45 Similarly, we merged all coral and associated habitats mapped in the Coral Reef
46 Habitat Map data to best match the protocol used for the Millennium data. Using
47 these two datasets combined, we calculated the total island reef habitat (<30 m depth)
48 within a 75 km buffer (~17,700 km²) at each site (*sensu* Mellin et al., 2010). We used
49 the Global Self-consistent, Hierarchical, High-resolution Geography Database

50 (<http://www.soest.hawaii.edu/pwessel/gshhg/>; Wessel & Smith, 1996) to estimate
51 land area (km²) within a 75 km buffer at each site.

52 Finally, we collated *in situ* estimates of structural complexity at each
53 stationary point count. CREP divers assessed structural complexity qualitatively on a
54 scale from 1 (very low) to 5 (very high) between 2010-2011, before switching method
55 and measuring maximum substrate height and proportion of area within substrate
56 height bins in each point count cylinder between 2012-2014 (Williams et al. 2015).
57 We merged both estimates by averaging complexity estimates across all sites for each
58 island that was sampled using both methods (n = 35), and then fitting the relationship
59 between qualitative complexity and maximum substrate height. We used this
60 relationship to convert qualitative complexity values to maximum substrate heights at
61 each 2010-11 site (Williams et al. 2015).

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63 **References**

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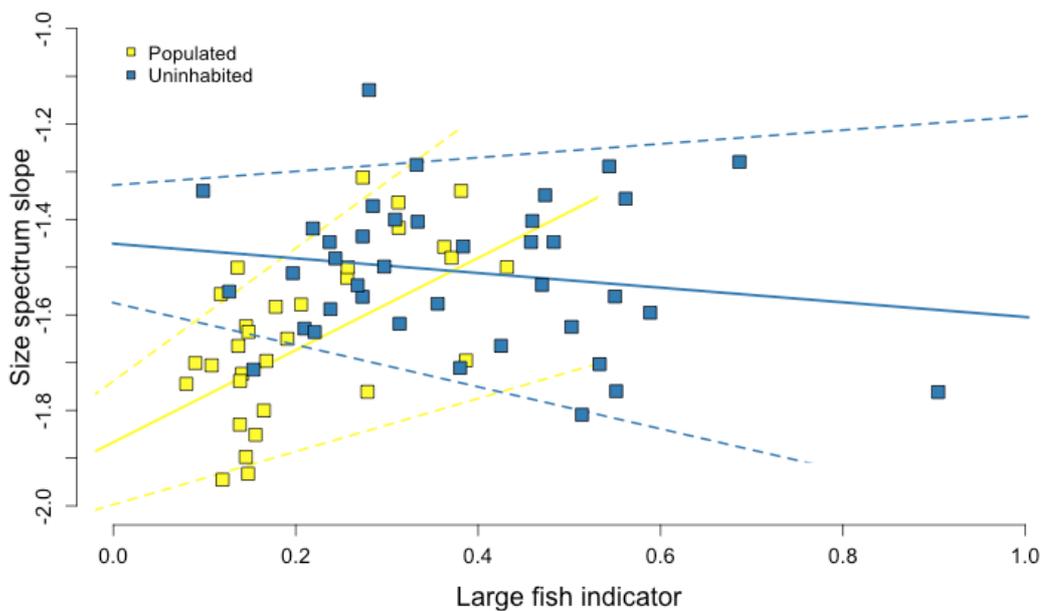
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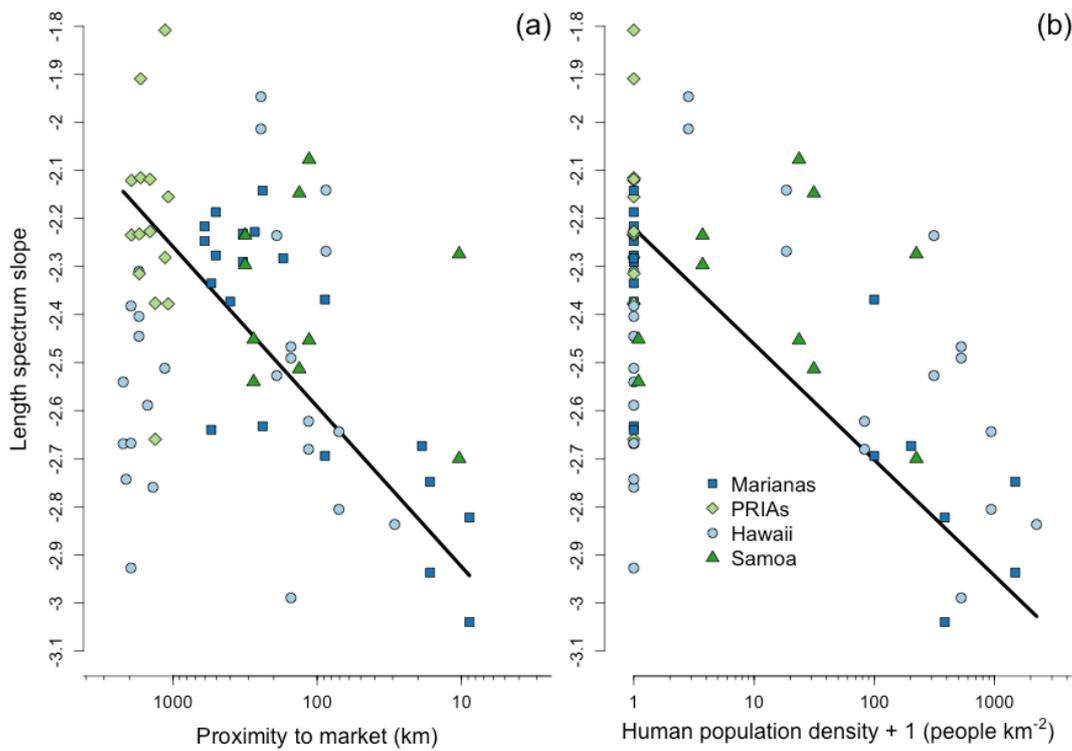
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108 Figure S1. Relationship between size spectra slopes and the LFI at populated (yellow)
109 and uninhabited (blue) reef areas. Linear regression (solid line) and 95% confidence
110 intervals (dashed lines) are parameter estimates from linear mixed effects model with
111 survey year as a random effect.

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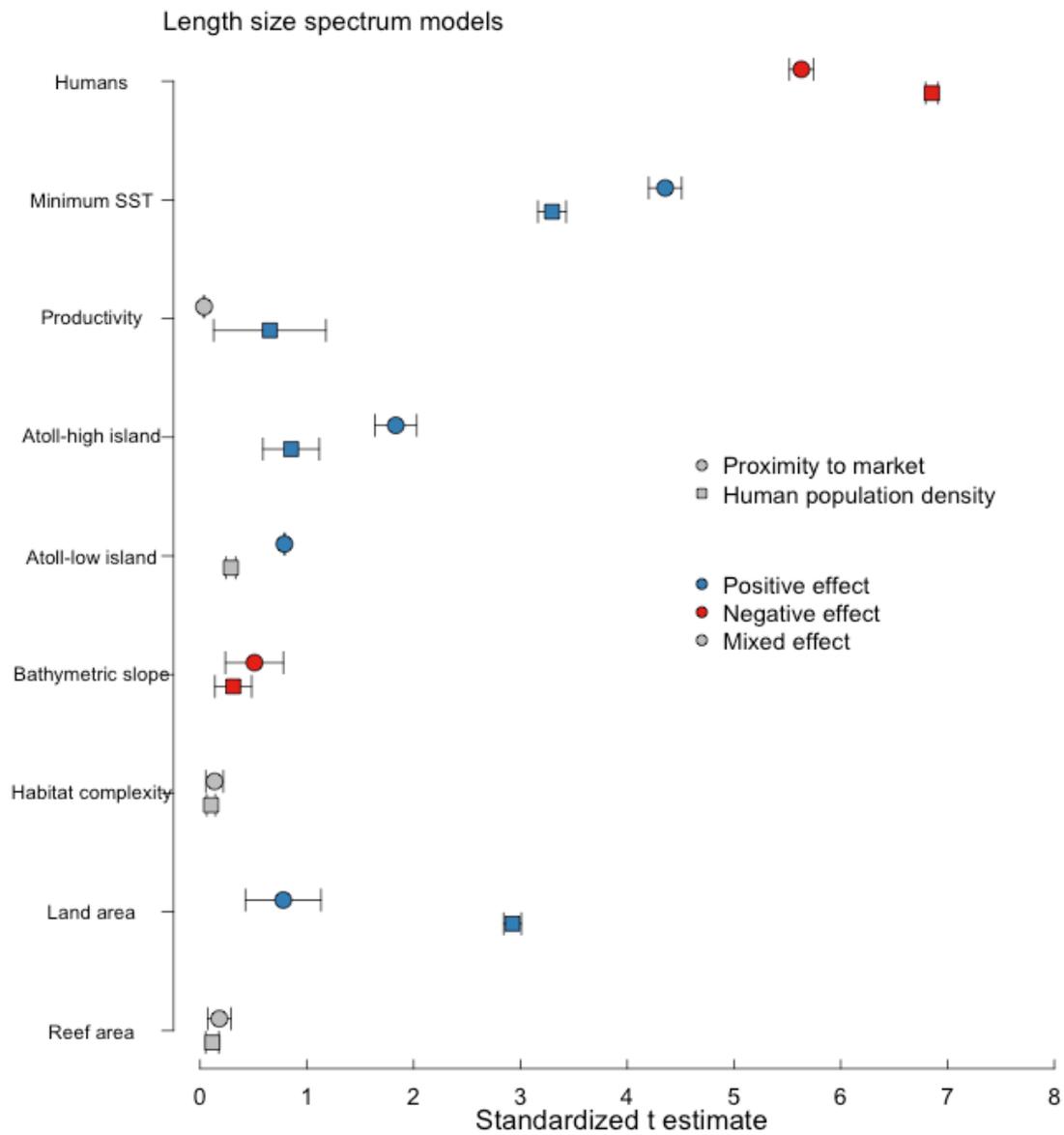
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119 Figure S2. Length spectra slopes across proximity to market (a) and human
120 population density (b). Length spectra relationships are model-averaged predictions
121 across the standardized range of observed log proximity to provincial capital (km) and
122 log human population density per forereef area (km²) (a, b respectively). Predictions
123 were made across the top model set ($\Delta AICc < 7$) and weighted using model
124 probabilities, while holding all other relevant covariates to their mean observed value
125 (Table S5). Dashed lines are the weighted sample variance at each value of human
126 covariate (though these are indistinguishable here from model predictions). For
127 visualization purposes, we included the observed data as points and coloured by
128 region (dark blue squares = Marianas archipelago; light blue circles = Hawaiian

129 archipelago, light green diamonds = Pacific Remote Island Areas, dark green triangles
 130 = American Samoa).



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 132 Figure S3. Model parameter estimates for length spectra. Length spectra are for the
 133 distance to market (circles) and human population density (squares) full model sets.
 134 Points are the weighted absolute t-values for each explanatory covariate, with
 135 weighted sample variance as error bars. T-values indicate the magnitude of each
 136 covariate effect, and colors indicate the direction of each covariate effect (blue =
 137 positive; red = negative; grey = mixed). See Table S5 for further details.