

Electronic Supplementary Material (ESM)

Community-wide scan identifies fish species associated with coral reef services across the Indo-Pacific

Eva Maire^{1,2,*}, Sébastien Villéger¹, Nicholas A. J. Graham³, Andrew S. Hoey², Joshua Cinner², Sebastian C. A. Ferse⁴, Catherine Aliaume¹, David J. Booth⁵, David A. Feary⁶, Michel Kulbicki⁷, Stuart A. Sandin⁸, Laurent Vigliola⁹ & David Mouillot^{1,2}

Additional information

Scales of data

Our data were organized at three spatial scales: reef site (n=1824), reef cluster (n=675), and nation/state (n=26).

- i) Reef site (the smallest scale, which had an average of 2.6 surveys/transects - hereafter 'reef').
- ii) Reef cluster (which had an average of 2.7 +/- 2.6 reef sites). Reefs within 4km of each other were clustered, and we used the centroid to estimate reef cluster-level socio-economic variables as described by Cinner *et al.* (2016) [1].
- iii) Nation/state (nation, state, or territory, which had an average of 26 +/- 42 reef clusters). A larger scale in our analysis was 'nation/state', which are jurisdictions that generally correspond to individual nations (but could also include states, territories, overseas regions), within which reef sites and reef clusters were nested for analysis.

Coral reef services

We considered fish biomass and live coral cover as two proxies for coral reef services as support, among many others, food security, shoreline protection and recreational value [2-5]. Fish biomass and coral cover are already monitored at the global scale using visual censuses as well as underwater video surveys (e.g. Reef Life Survey, Catlin Sea Survey) and are highly sensitive to local human activities (e.g. fishing, habitat destruction, pollution) and global climate change [6] and thus, can be considered as key variables for the health and productivity of coral reefs [1, 7, 8].

Fish Biomass

Reef fish biomass estimates were based on instantaneous visual counts from 4,694 surveys collected from 1,824 reefs (Fig. S1). Surveys were carried out using two census methods (belt-transects or distance sampling) and were conducted between 2004 and 2013. On average 2.4 transects (sd=1.32; range: 1-10) were performed on each reef. Within each surveyed area, reef associated fishes were identified to species level, abundance was counted, and total length (TL) estimated.

To make estimates of biomass from these transect-level data comparable among studies, we:

- i) Considered only Indo-Pacific reefs and retained families that were consistently included in surveys and were above a minimum size cut-off. Thus, we only retained counts of non-cryptic reef fish species >10cm in total length, that are reef-associated (30 families, 748 species) (Table S1). We did not include sharks as they were often excluded from visual surveys. We calculated total biomass of fishes on each reef using published species-level length-weight relationship parameters or those available on FishBase [39]. When length-weight relationship parameters were not available for a species, we used the parameters for a closely related species or genus.
- ii) Depth and habitat were included as environmental variables in the model.
- iii) Reef fish biomass estimates were based on instantaneous visual counts using belt-transects or distance sampling. Because several biases were associated with these two methods [9] and the total area of transects changed between locations, we included census method and sampling area as covariates in the model.

Biomass values were calculated at the reef scale and showed a high variability (mean=1,055 kg.ha⁻¹; range: 2-25,910 kg.ha⁻¹).

Live coral cover

Percent cover of live coral was based on 1,715 point-intercept transects or quadrats collected from 741 reefs (Fig. S2). All surveys were conducted between 2008 and 2013. On average 2.3 transects (sd=1.03; range: 1-4) were performed on each reef. To make estimates of coral cover from these transect-level data comparable among studies, we included depth, habitat and census method (which already different in the sampling area) as covariates in the model (see details in Supplementary Material).

Coral cover values were calculated at the reef scale and showed a high variability (mean=27%; range: 3-94%).

Socioeconomic variables

-Gravity of Human Populations: We used two gravity indices (the cumulative gravity of all cities within 500km and the gravity of the nearest human population) for each of our reefs where we had in situ ecological data because those had the strongest relationships with reef fish biomass as described in Cinner *et al.* (2016) [1].

-Management: For each reef, we determined if it was: i) marine reserve- whether it fell within the borders of a no-take marine reserve and we asked data providers to further classify whether the reserve had high or low levels of compliance; ii) restricted - whether there were active restrictions on gears (e.g. bans on the use of nets, spearguns, or traps) or fishing effort (which could have included areas inside marine protected areas that were not necessarily no take); or iii) fished - regularly fished without effective restrictions. To determine these classifications, we used the expert opinion of the data providers, and validated this with a global database of marine reserve boundaries [10].

-Local Human Population Growth: We created a 100km buffer around each reef cluster and used this to calculate human population within the buffer in 2000 and 2010 based on the Socioeconomic Data and Application Centre (SEDAC) gridded population of the world database. Population growth was the proportional difference between the population in 2000 and 2010. We chose a 100km buffer as a reasonable range at which many key human impacts from population (e.g., land-use and nutrients) might affect reefs [11].

-Human Development Index (HDI): HDI is a summary measure of human development encompassing: a long and healthy life, being knowledgeable, and having a decent standard of living. In cases where HDI values were not available specific to the State (e.g. Florida and Hawaii), we used the national (e.g. USA) HDI value.

-Population Size: For each nation/state, we determined the size of the human population. Data were derived mainly from census reports, the CIA fact book, and Wikipedia.

-Tourism: We examined tourist arrivals relative to the nation/state population size (above). Tourism arrivals were gathered primarily from the World Tourism Organization's Compendium of Tourism Statistics.

-National Reef Fish Landings: Catch data were obtained from the Sea Around Us Project (SAUP) catch database (www.seaaroundus.org), except for Florida, which was not reported separately in the database. We identified 200 reef fish species and taxon groups in the SAUP catch database to define landings in tons/km² of reef.

-Voice and Accountability: This metric, from the World Bank survey on governance, reflects the perceptions of the extent to which a country's citizens are able to participate in selecting their government, as well as freedom of expression, freedom of association, and a free media. In cases where governance values were not available specific to the Nation/state (e.g. Florida and Hawaii), we used national (e.g. USA) values.

Environmental variables

-Depth: The depth of reef surveys was grouped into the following categories: <4m, 4-10m, >10m to account for broad differences in reef fish community structure attributable to a number of inter-linked depth-related factors. Categories were necessary to standardize methods used by data providers and were determined by pre-existing categories used by several data providers.

-Habitat: We included the following habitat categories: i) Slope: The reef slope habitat is typically on the ocean side of a reef, where the reef slopes down into deeper water; ii) Crest: The reef crest habitat is the section that joins a reef slope to the reef flat. The zone is typified by high wave energy (i.e. where the waves break) and a change in the angle of the reef from an inclined slope to a horizontal reef flat; iii) Flat: The reef flat habitat is typically horizontal and extends back from the reef crest for 10's to 100's of meters; iv) Lagoon / back reef: Lagoonal reef habitats are where the continuous reef flat breaks up into more patchy reef environments sheltered from wave energy. These habitats can be behind barrier / fringing reefs or within atolls. Back reef habitats are similar broken habitats where the wave energy does not typically reach the reefs and thus forms a less continuous 'lagoon style' reef habitat.

For this analysis, we excluded other less prevalent habitat types (channels and banks) and we verified the sites' habitat information using Google Earth, and site depth information.

-Productivity: We extracted oceanic productivity for each reef following the procedure described by Gove et al. (2013) [12]. We delimited a 100km buffer around each site, we removed shallow waters pixels (those that intersected or were contained within the depth contour of 30m from the General Bathymetric Chart of the Oceans 2014

(<http://www.gebco.net/>), a global gridded bathymetry dataset) and then calculated the average of monthly chlorophyll-a concentration (proxy for phytoplankton biomass) using data provided at a 4km-resolution by Aqua MODIS (Moderate Resolution Imaging Spectro-radiometer) for years 2005 to 2010 (in hdf format).

Analyses

We used linear mixed models to analyze biomass and live coral cover after checking that both log-transformed variables were normally distributed. For both models, we set site, regional locations and nation/state as random effects to account for the hierarchical nature of the data (i.e. reefs nested in sites, sites nested in regional locations and in nations/states). All continuous covariates were standardized for the analysis and their corresponding Akaike weights were computed to assess their importance (Table S2 & S3).

To check the fit of the linear mixed model, we checked for the representation of actual versus predicted values and we calculated the accuracy of the two models which came to 88% for biomass and 83% for coral cover. To examine homoscedasticity, we checked residuals against fitted values. We checked that the residuals were normally distributed. All analyses were undertaken using R (3.3) statistical packages.

Functional traits to describe fish species.

The 381 candidate fish species were functionally described using six traits extracted from Mouillot et al. (2014) [13]: (1) size, coded using 3 ordered categories: 10-30cm, 30.1-50cm, >50cm ; (2) mobility, coded using 3 ordered categories: sedentary, mobile within a reef and mobile between reefs; (3) period of activity, coded using 3 ordered categories: diurnal, both diurnal and nocturnal, and nocturnal ; (4) schooling, coded using 5 ordered categories: solitary, pairing, or living in small (3-20 individuals), medium (20-50 individuals) or large groups (>50 groups) ; (5) vertical position in the water column, coded using 3 ordered categories: benthic, benthic-pelagic and pelagic; (6) diet, coded using 7 trophic categories: herbivorous-detrivorous, macro-algal browser, invertivorous targeting sessile invertebrates, invertivorous targeting mobile invertebrates, planktivorous, piscivorous, and omnivorous, i.e. fish for which both plant and animal material are important in their diet. Since all traits were categorical, species with identical traits were grouped into functional entities.

Defining presence of species based on abundance threshold.

The community-wide scan (CWS) approach can be adapted for a wide range of taxa from all the ecosystems. The way candidates are tested can be modulated while respecting independence between predicted and explanatory variables (binary variables are a convenient way to test the effect of candidates). More specifically, presence of terrestrial or marine taxa can be determined using any convenient abundance threshold such as a minimum number of individuals, cover rate or biomass.

As an application, we tested two procedures to define presence of reef fish species based on a biomass threshold and compared key species found with those two procedures:

i) the relative intraspecific biomass which defines presence of a species in a community as soon as its biomass reaches upper percentiles of the distribution of biomass of this species over all studied communities. We used the 99th and 95th percentiles (the top 1 and 5%) of the biomass distribution over reef sites as thresholds. This approach is particularly relevant for species widely distributed with normal distribution of biomass while it may not be used for species infrequently encountered or with a skewed biomass distribution.

ii) the relative community biomass which defines presence as soon as the focal species reach a defined minimum percentage of the total biomass of the fish community. Hence, this approach is not affected by biomass distribution among sites. As two thresholds, we tested a contribution of species to total biomass in excess of 1 and 5%, respectively.

Figures and Tables

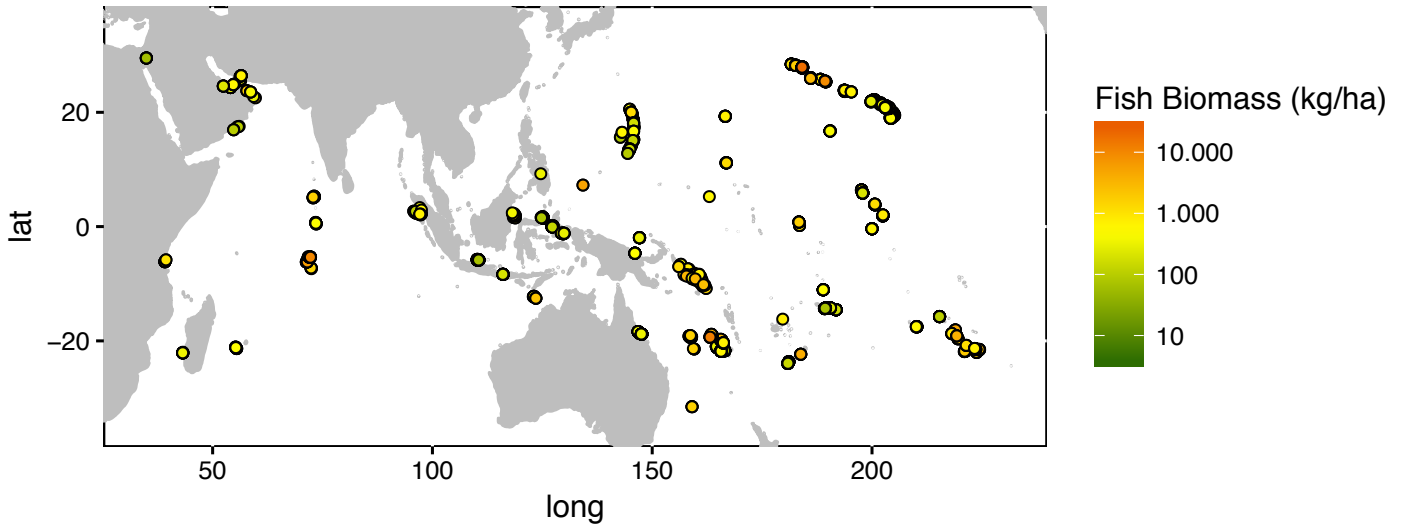


Figure S1. Map of the reef fish biomass observed in 1,824 reefs located in the Indo-Pacific.

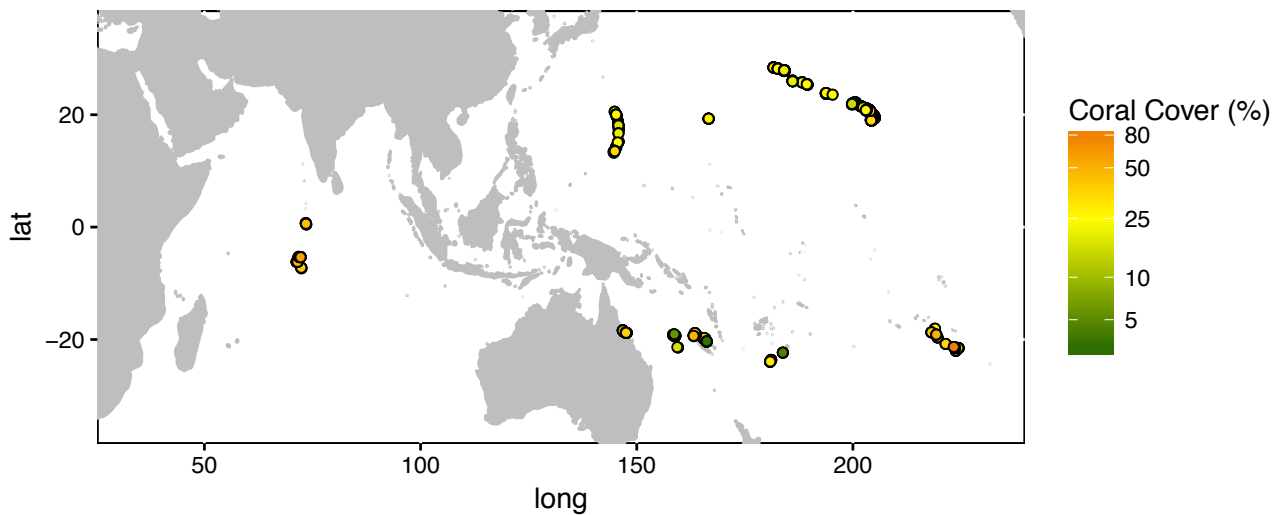


Figure S2. Map of the live coral cover observed in 741 reefs located in the Indo-Pacific. Information on coral cover was not available for all sites at which reef fish biomass was surveyed.

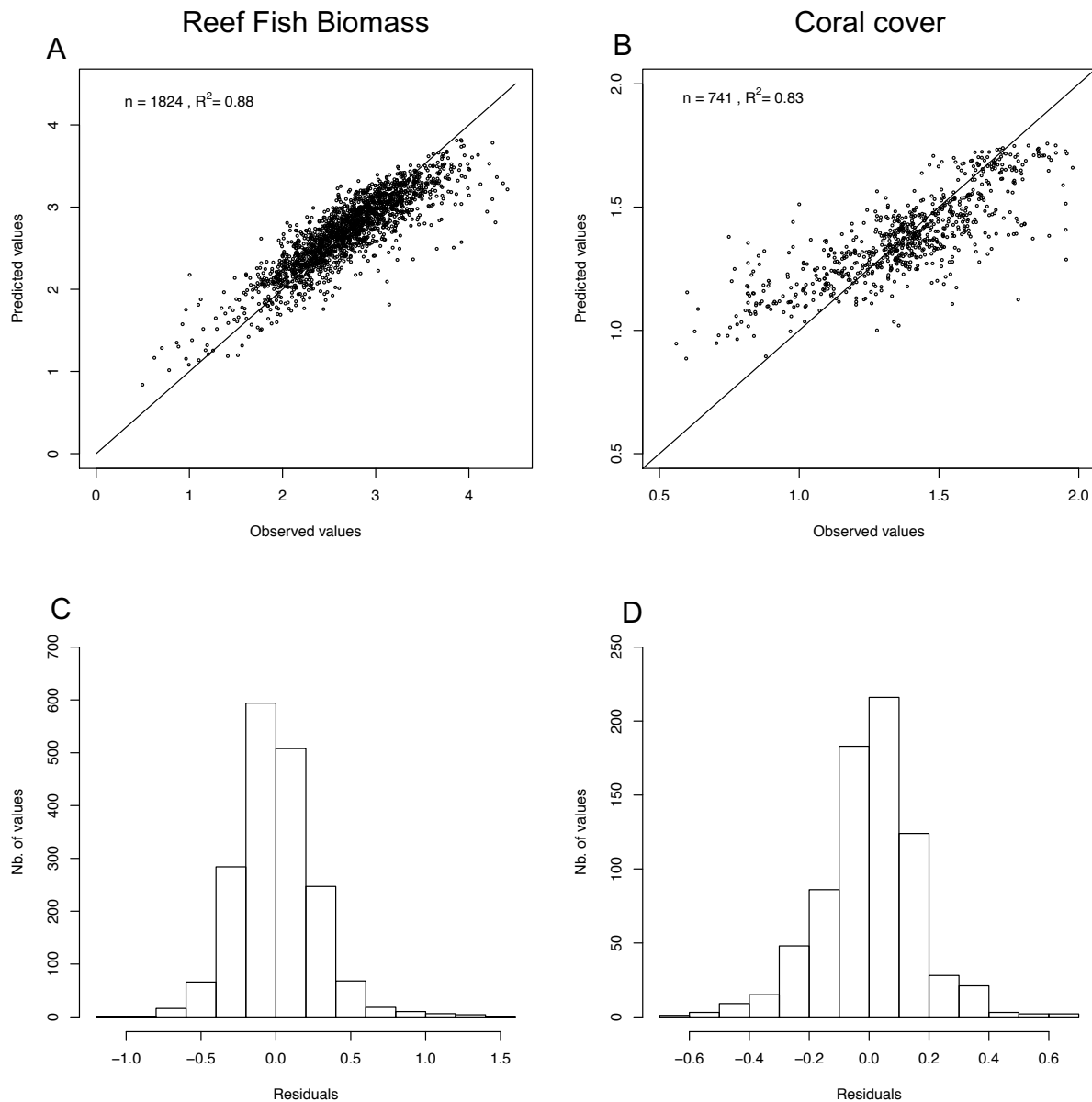


Figure S3. Accuracy and residuals of the two initial models (fish biomass and coral cover).

The accuracy of each model is assessed using the Pearson correlation between predicted and observed values which came to 88% for biomass (A) and 83% for coral cover (B). We checked that residuals of the initial model of fish biomass (C) and coral cover (D) were normally distributed.

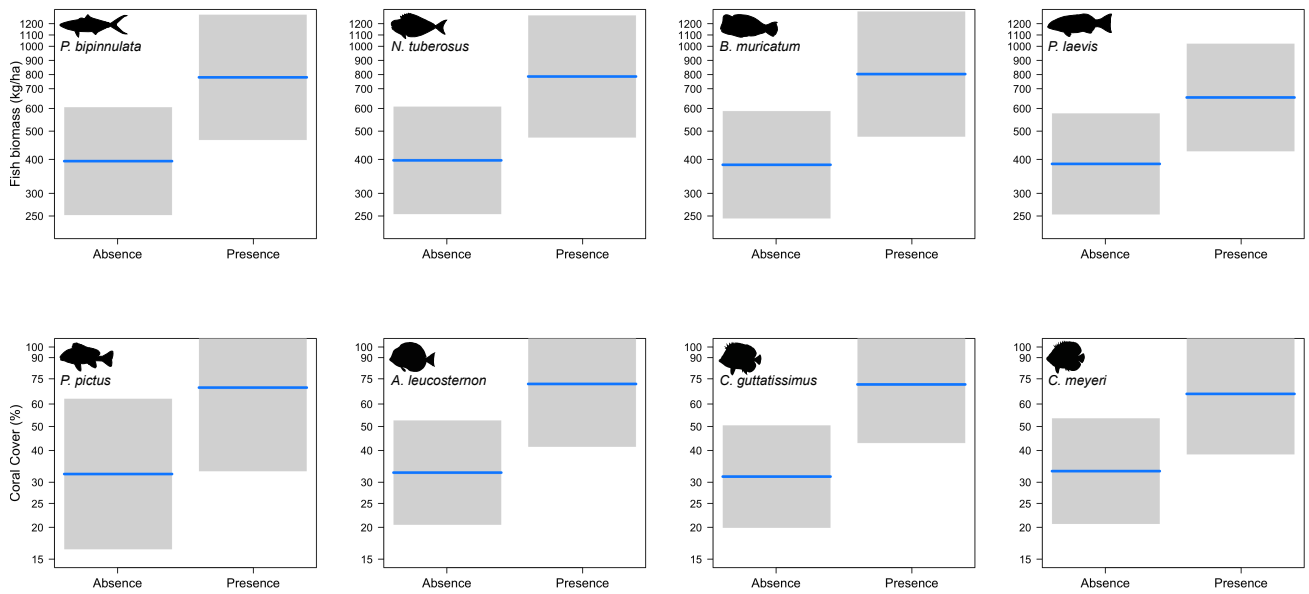


Figure S4. Net effect of the 4 key fish species linked to the highest levels of fish biomass (top) and live coral cover (bottom). We extracted the pure effects of all the key species and determined which were related to the highest levels of biomass and coral cover respectively, using a partial plot from each LMM while the other variables are held constant.

Table S1: List of coral reef fish families included in the study and their common name.

Families included are: Acanthuridae, Balistidae, Caesionidae, Carangidae, Chaetodontidae, Cirrhitidae, Diodontidae, Ehippidae, Haemulidae, Holocentridae, Kyphosidae, Labridae, Lethrinidae, Lutjanidae, Monacanthidae, Mullidae, Nemipteridae, Pempheridae, Pinguipedidae, Pomacanthidae, Priacanthidae, Pseudochromidae, Scombridae, Serranidae, Siganidae, Sparidae, Sphyraenidae, Synodontidae, Tetradontidae, Zanclidae.

Fish family	Common family name	Fish family	Common family name
Acanthuridae	Surgeonfishes	Mullidae	Goatfishes
Balistidae	Triggerfishes	Nemipteridae	Coral Breams
Caesionidae	Fusiliers	Pempheridae	Sweepers
Carangidae	Jacks/Trevallies	Pinguipedidae	Sandperches
Chaetodontidae	Butterflyfishes	Pomacanthidae	Angelfishes
Cirrhitidae	Hawkfishes	Priacanthidae	Bigeyes
Diodontidae	Porcupinefishes	Pseudochromidae	Dottybacks
Ehippidae	Batfishes	Scombridae	Mackerels and Tunas
Haemulidae	Sweetlips	Serranidae	Groupers
Holocentridae	Squirrelfishes, Soldierfishes	Siganidae	Rabbitfishes
Kyphosidae	Drummers	Sparidae	Porgies
Labridae	Wrasses and Parrotfishes	Sphyraenidae	Barracuda
Lethrinidae	Emperors	Synodontidae	Lizardfishes
Lutjanidae	Snappers	Tetradontidae	Puffers
Monacanthidae	Filefishes	Zanclidae	Moorish Idol

Table S2. Relative importance of variables of the fish biomass initial model. The relative importance of each variable is based on the sum of Akaike weights of all the possible models (i.e. all the possible combination of variables) in which the variable is present.

Relative variable importance	
Biodiversity	
Species richness	1
Environmental variables	
Depth	1
Oceanic productivity	1
Habitat	0.94
Socio-economic variables	
Population size	1
Management	1
Tourism	1
HDI	0.72
Reef fish landings	0.71
Gravity of markets	0.66
Voice accountability	0.62
Local population growth	0.39
Gravity of human settlement	0.28
Method	
Sampling area	1
Census method	1

Table S3. Relative importance of variables of the coral cover initial model. The relative importance of each variable is based on the sum of Akaike weights of all the possible models (i.e. all the possible combination of variables) in which the variable is present.

<i>Relative variable importance</i>	
<i>Biodiversity</i>	
Species richness	1
<i>Environmental variables</i>	
Oceanic productivity	1
Habitat	1
Depth	0.56
<i>Socio-economic variables</i>	
Population size	1
Tourism	1
Local population growth	0.85
HDI	0.74
Gravity of markets	0.48
Reef fish landings	0.44
Management	0.41
Voice accountability	0.37
Gravity of human settlement	0.27
<i>Method</i>	
Census method	1

Table S4. Details of key fish species for reef fish biomass. AIC of the initial model M_0 (environment, socio-economics and species richness) provides reference Akaike Information Criterion (AIC M_0). Presence of each species has been added to M_0 as binary variable using occurrence (i.e. presence of at least 1 individual). For each species, number of occurrences, coefficient in the model (Coeff.) and AIC of the model accounting for the species (AIC M_1) are reported. Comparisons with reference AIC (AIC M_0) are also provided (Δ AIC).

Fish species	Nb of occ.	Coeff.	AIC M_1	Δ AIC	AIC M_0
<i>Caranx ignobilis</i>	98	0.29	1166.6	54.6	1221.2
<i>Lutjanus bohar</i>	448	0.18	1171.4	49.8	1221.2
<i>Plectropomus laevis</i>	129	0.23	1185.7	35.5	1221.2
<i>Aprion virescens</i>	169	0.18	1189.7	31.5	1221.2
<i>Macolor niger</i>	157	0.2	1189.9	31.3	1221.2
<i>Chlorurus microrhinos</i>	314	0.15	1190	31.2	1221.2
<i>Hipposcarus longiceps</i>	163	0.19	1193	28.2	1221.2
<i>Naso tuberosus</i>	51	0.3	1195.9	25.3	1221.2
<i>Lethrinus atkinsoni</i>	74	0.28	1198.3	22.9	1221.2
<i>Bolbometopon muricatum</i>	38	0.32	1199.9	21.3	1221.2
<i>Cheilinus undulatus</i>	134	0.17	1200	21.2	1221.2
<i>Elagatis bipinnulata</i>	32	0.3	1201.5	19.7	1221.2
<i>Naso hexacanthus</i>	113	0.16	1203.1	18.1	1221.2
<i>Caranx melampygus</i>	286	0.11	1207.1	14.1	1221.2
<i>Scarus altipinnis</i>	181	0.13	1208.5	12.7	1221.2
<i>Naso caesioides</i>	31	0.27	1209.2	12	1221.2
<i>Parupeneus crassilabris</i>	55	0.2	1212.3	8.9	1221.2
<i>Naso unicornis</i>	380	0.09	1212.3	8.9	1221.2
<i>Lethrinus olivaceus</i>	79	0.16	1212.8	8.4	1221.2
<i>Macolor macularis</i>	107	0.16	1214.7	6.5	1221.2
<i>Lutjanus gibbus</i>	191	0.12	1215	6.2	1221.2
<i>Gymnosarda unicolor</i>	23	0.24	1215.1	6.1	1221.2
<i>Acanthurus albipectoralis</i>	26	0.23	1215.7	5.5	1221.2
<i>Acanthurus lineatus</i>	282	0.09	1215.8	5.4	1221.2
<i>Scarus rubroviolaceus</i>	367	0.08	1216.6	4.6	1221.2
<i>Acanthurus dussumieri</i>	168	0.1	1217	4.2	1221.2

Table S5. Details of key fish species for live coral cover. AIC of the initial model M_0 (environment, socio-economics and species richness) provides reference Akaike Information Criterion (AIC M_0). Presence of each species has been added to M_0 as binary variable using occurrence (i.e. presence of at least 1 individual). For each species, number of occurrences, coefficient in the model (Coeff.) and AIC of the model accounting for the species (AIC M_1) are reported. Comparisons with reference AIC (AIC M_0) are also provided (Δ AIC).

Fish species	Nb of occ.	Coeff.	AIC M_1	Δ AIC	AIC M_0
<i>Chaetodon trifascialis</i>	132	0.22	-106.6	66.4	-40.2
<i>Hemigymnus fasciatus</i>	166	0.17	-82.1	41.9	-40.2
<i>Bodianus loxozonus</i>	35	0.27	-69.1	28.9	-40.2
<i>Zebrasoma scopas</i>	101	0.16	-66.1	25.9	-40.2
<i>Chaetodon guttatissimus</i>	40	0.35	-60.5	20.3	-40.2
<i>Scarus frenatus</i>	158	0.15	-59.1	18.9	-40.2
<i>Chaetodon baronessa</i>	65	0.22	-58.3	18.1	-40.2
<i>Plectorhinchus picus</i>	10	0.33	-55.1	14.9	-40.2
<i>Oxycheilinus unifasciatus</i>	147	0.1	-54.1	13.9	-40.2
<i>Chaetodon meyeri</i>	40	0.29	-54	13.8	-40.2
<i>Chlorurus strongylocephalus</i>	32	0.26	-53.9	13.7	-40.2
<i>Lutjanus gibbus</i>	55	0.16	-53.6	13.4	-40.2
<i>Chaetodon unimaculatus</i>	48	0.16	-52.8	12.6	-40.2
<i>Chaetodon pelewensis</i>	70	0.16	-52.7	12.5	-40.2
<i>Acanthurus leucosternon</i>	44	0.33	-52.6	12.4	-40.2
<i>Ctenochaetus truncatus</i>	36	0.27	-50.6	10.4	-40.2
<i>Chaetodon reticulatus</i>	86	0.18	-50.5	10.3	-40.2
<i>Acanthurus albipectoralis</i>	24	0.2	-49.9	9.7	-40.2
<i>Labrichthys unilineatus</i>	59	0.16	-49.4	9.2	-40.2
<i>Lutjanus bohar</i>	199	0.11	-48.9	8.7	-40.2
<i>Bodianus axillaris</i>	118	0.1	-47.9	7.7	-40.2
<i>Acanthurus thompsoni</i>	88	0.11	-47.3	7.1	-40.2
<i>Scarus chameleon</i>	74	0.11	-47	6.8	-40.2
<i>Paracirrhites nesus</i>	24	0.22	-46.5	6.3	-40.2
<i>Scarus tricolor</i>	31	0.2	-46.2	6	-40.2
<i>Cephalopholis urodeta</i>	145	0.13	-45.2	5	-40.2
<i>Scarus globiceps</i>	76	0.12	-44.7	4.5	-40.2
<i>Cantherhines dumerilii</i>	52	0.11	-44.7	4.5	-40.2

Table S6. Functional traits of key fish species for reef fish biomass and live coral cover.

23, 25 and 3 fish species are significantly related to biomass (BM), live coral cover (CC) or both (BOTH) of those proxies of coral reef services respectively. Each species is described using six traits: (1) size, coded using 3 ordered categories: 10-30cm, 30.1-50cm, >50cm; (2) mobility, coded using 3 ordered categories: sedentary, mobile within a reef and mobile between reefs; (3) period of activity, coded using 3 ordered categories: diurnal, both diurnal and nocturnal, and nocturnal; (4) schooling, coded using 5 ordered categories: solitary, pairing, or living in small (3-20 individuals), medium (20-50 individuals) or large groups (>50 groups); (5) vertical position in the water column, coded using 3 ordered categories: benthic, benthopelagic and pelagic; (6) diet, coded using 7 trophic categories: herbivorous-detritivorous (HD), macro-algal herbivorous (HM), invertivorous targeting sessile invertebrates (IS), invertivorous targeting mobile invertebrates (IM), planktivorous (PK), piscivorous (FC), and omnivorous (OM), i.e. fish for which both vegetal and animal material are important in their diet. Several species can have the same functional traits (trait combinations with the same Funct. Entity ID) and thus, represent the same functional entity. In total, 51 key fish species are described, which represent 35 different functional entities with 6 entities common to both fish biomass and coral cover (highlighted in grey).

ES proxy	Fish Species	Size Class	Mobility	Activity	Schooling	Position	Diet	Funct. Entity ID
BM	<i>Acanthurus dussumieri</i>	10-30cm	Mob	Day	SmallG	Bottom	HD	6
BM	<i>Acanthurus lineatus</i>	10-30cm	Sed	Day	Sol	Bottom	HD	15
BM	<i>Aprion virescens</i>	50-153cm	VMob	Both	SmallG	Low	FC	33
BM	<i>Bolbometopon muricatum</i>	50-153cm	VMob	Day	MedG	Bottom	IS	35
BM	<i>Caranx ignobilis</i>	50-153cm	VMob	Both	Sol	High	FC	34
BM	<i>Caranx melampygus</i>	30-50cm	VMob	Both	SmallG	High	FC	25
BM	<i>Cheilinus undulatus</i>	50-153cm	Mob	Day	Sol	Bottom	IM	31
BM	<i>Chlorurus microrhinos</i>	30-50cm	Mob	Day	SmallG	Bottom	OM	23
BM	<i>Elagatis bipinnulata</i>	50-153cm	VMob	Both	MedG	High	FC	32
BM	<i>Gymnosarda unicolor</i>	50-153cm	VMob	Both	Sol	High	FC	34
BM	<i>Hipposcarus longiceps</i>	10-30cm	Mob	Day	MedG	Bottom	OM	3
BM	<i>Lethrinus atkinsoni</i>	10-30cm	VMob	Night	MedG	Bottom	IM	20
BM	<i>Lethrinus olivaceus</i>	30-50cm	VMob	Both	Sol	Bottom	FC	26
BM	<i>Macolor macularis</i>	10-30cm	Mob	Both	MedG	High	PK	1
BM	<i>Macolor niger</i>	30-50cm	Mob	Both	MedG	High	PK	22
BM	<i>Naso caesius</i>	30-50cm	VMob	Day	MedG	High	PK	29
BM	<i>Naso hexacanthus</i>	30-50cm	VMob	Day	LargeG	High	PK	27
BM	<i>Naso tuberosus</i>	30-50cm	VMob	Day	MedG	High	HM	28
BM	<i>Naso unicornis</i>	10-30cm	Mob	Day	SmallG	Bottom	HM	7
BM	<i>Parupeneus crassilabris</i>	10-30cm	Mob	Day	SmallG	Bottom	IM	8
BM	<i>Plectropomus laevis</i>	50-153cm	Mob	Day	Sol	Bottom	FC	30
BM	<i>Scarus altipinnis</i>	10-30cm	Mob	Day	LargeG	Bottom	OM	2
BM	<i>Scarus rubroviolaceus</i>	30-50cm	Mob	Day	SmallG	Bottom	OM	23
CC	<i>Acanthurus leucosternon</i>	10-30cm	VMob	Day	LargeG	Bottom	HD	18
CC	<i>Acanthurus thompsoni</i>	10-30cm	VMob	Day	MedG	Low	PK	19
CC	<i>Bodianus axillaris</i>	10-30cm	Mob	Day	Sol	Bottom	IM	9
CC	<i>Bodianus loxozonus</i>	10-30cm	Mob	Day	Sol	Bottom	IM	9
CC	<i>Cantherhines dumerilii</i>	10-30cm	Mob	Day	Pair	Bottom	IS	4
CC	<i>Cephalopholis urodeta</i>	10-30cm	Sed	Both	Sol	Bottom	FC	11
CC	<i>Chaetodon baronessa</i>	10-30cm	Sed	Day	Pair	Bottom	IS	12
CC	<i>Chaetodon guttattissimus</i>	10-30cm	Sed	Day	Pair	Bottom	IS	12
CC	<i>Chaetodon meyeri</i>	10-30cm	Sed	Day	Pair	Bottom	IS	12
CC	<i>Chaetodon pelewensis</i>	10-30cm	Sed	Day	Pair	Bottom	IS	12
CC	<i>Chaetodon reticulatus</i>	10-30cm	Sed	Day	Pair	Low	IS	13
CC	<i>Chaetodon trifascialis</i>	10-30cm	Sed	Day	Pair	Bottom	IS	12
CC	<i>Chaetodon unimaculatus</i>	10-30cm	Sed	Day	Pair	Bottom	IS	12
CC	<i>Chlorurus strongylocephalus</i>	30-50cm	Mob	Day	SmallG	Bottom	OM	23
CC	<i>Ctenochaetus truncatus</i>	10-30cm	Sed	Day	SmallG	Bottom	OM	14
CC	<i>Hemigymnus fasciatus</i>	10-30cm	Mob	Day	Sol	Bottom	IM	9
CC	<i>Labrichthys unilineatus</i>	10-30cm	Sed	Day	Sol	Bottom	IS	17
CC	<i>Oxycheilinus unifasciatus</i>	10-30cm	Mob	Day	SmallG	Bottom	FC	5
CC	<i>Paracirrhites nesus</i>	10-30cm	Sed	Day	Sol	Bottom	IM	16
CC	<i>Plectorhinchus picus</i>	50-153cm	Mob	Day	Sol	Bottom	IM	31
CC	<i>Scarus chameleon</i>	10-30cm	Mob	Day	Sol	Bottom	OM	10
CC	<i>Scarus frenatus</i>	10-30cm	Mob	Day	Sol	Bottom	OM	10
CC	<i>Scarus globiceps</i>	10-30cm	Mob	Day	Sol	Bottom	OM	10
CC	<i>Scarus tricolor</i>	10-30cm	Mob	Day	Sol	Bottom	OM	10
CC	<i>Zebrasoma scopas</i>	10-30cm	Mob	Day	SmallG	Bottom	HD	6
BOTH	<i>Acanthurus albipectoralis</i>	10-30cm	VMob	Day	MedG	Low	PK	19
BOTH	<i>Lutjanus bohar</i>	30-50cm	Mob	Night	MedG	Low	FC	24
BOTH	<i>Lutjanus gibbus</i>	30-50cm	Mob	Both	MedG	Bottom	IM	21

Table S7. Comparative results of potential key species for fish biomass using relative intra-community (left) and intra-specific (right) biomass to define presence (threshold defined as 1%). AIC of the initial model M_0 provides reference Akaike Information Criterion (AIC M_0). Presence of each species has been added to M_0 as binary variable using 2 criteria: i) contribution of species to total biomass higher than 1% and ii) the 99th percentile of the biomass distribution of each species. For each species, number of raw occurrences, number of occurrences meeting the biomass threshold, and AIC of the model accounting for the species (AIC M_1) are reported. For clarity, only key species are presented and those that are significantly related to fish biomass under the 2 approaches are highlighted in grey. One species had an AIC lower than M_0 but did not exceed the performance criterion ($\Delta AIC > 4$) to be considered as key species (underlined).

INITIAL MODEL (M_0): AIC = 1221.2							
INTRA-COMMUNITY				INTRA-SPECIFIC			
Fish species	Nb of occ.	Nb of occ. (1%)	AIC M_1	Fish species	Nb of occ.	Nb of occ. (1%)	AIC M_1
<i>Caranx ignobilis</i>	98	96	1166.6	<i>Caranx ignobilis</i>	98	19	1157.1
<i>Lutjanus bohar</i>	448	332	1185.0	<i>Lutjanus gibbus</i>	191	19	1177.5
<i>Plectropomus laevis</i>	129	116	1199.3	<i>Macolor niger</i>	157	19	1177.8
<i>Chlorurus microrhinos</i>	314	285	1202.3	<i>Naso hexacanthus</i>	113	19	1183.8
<i>Bolbometopon muricatum</i>	38	35	1206.1	<i>Lethrinus atkinsoni</i>	74	19	1188.2
<i>Naso tuberosus</i>	51	46	1206.6	<i>Bolbometopon muricatum</i>	38	20	1188.4
<i>Macolor niger</i>	157	100	1209.8	<i>Hipposcarus longiceps</i>	163	19	1191.6
<i>Aprion virescens</i>	169	151	1210.9	<i>Scarus altipinnis</i>	181	19	1192.8
<i>Hipposcarus longiceps</i>	163	107	1211.5	<i>Chlorurus microrhinos</i>	314	20	1194
<i>Naso hexacanthus</i>	113	86	1212.6	<i>Lutjanus kasmira</i>	202	19	1196
<i>Lethrinus atkinsoni</i>	74	58	1213.3	<i>Lutjanus bohar</i>	448	19	1197.3
<i>Elagatis bipinnulata</i>	32	28	1214.6	<i>Plectropomus laevis</i>	129	19	1202.2
<i>Naso caesius</i>	31	22	1216.7	<i>Naso tuberosus</i>	51	20	1202.7
				<i>Acanthurus blochii</i>	225	19	1202.9
				<i>Siganus punctatus</i>	50	19	1203.6
				<i>Naso unicornis</i>	380	19	1203.7
				<i>Naso caesius</i>	31	19	1204.1
				<i>Ctenochaetus striatus</i>	954	19	1205.1
				<i>Caranx melampygus</i>	286	19	1205.3
				<i>Chlorurus bleekeri</i>	210	19	1206.1
				<i>Macolor macularis</i>	107	19	1206.5
				<i>Caesio cuning</i>	110	19	1206.7
				<i>Chlorurus frontalis</i>	75	19	1206.9
				<i>Monotaxis grandoculis</i>	488	19	1206.9
				<i>Scarus ghobban</i>	210	19	1208.7
				<i>Pterocaesio tile</i>	181	19	1208.9
				<i>Acanthurus lineatus</i>	282	19	1208.9
				<i>Plectorhynchus orientalis</i>	33	19	1209.4
				<i>Elagatis bipinnulata</i>	32	19	1209.9
				<i>Naso vlamingii</i>	51	19	1211
				<i>Scolopsis ghanam</i>	35	19	1211.2
				<i>Lutjanus fulviflamma</i>	107	20	1212.2
				<i>Acanthurus xanthopterus</i>	57	20	1212.2
				<i>Cheilinus undulatus</i>	134	19	1212.4
				<i>Naso brevirostris</i>	155	19	1213.6
				<i>Naso lituratus</i>	624	19	1213.6
				<i>Parupeneus barberinus</i>	55	20	1213.8
				<i>Lethrinus olivaceus</i>	79	23	1213.9
				<i>Caesio teres</i>	94	19	1214
				<i>Balistoides viridescens</i>	82	19	1214.6
				<i>Cetoscarus bicolor</i>	144	19	1214.6
				<i>Pterocaesio marri</i>	28	19	1214.9
				<i>Scarus rubroviolaceus</i>	367	19	1215.2
				<i>Kyphosus vaigiensis</i>	52	19	1215.4
				<i>Scarus rivulatus</i>	176	19	1215.5
				<i>Epibulus insidiator</i>	428	19	1215.7
				<i>Cetoscarus ocellatus</i>	51	19	1215.7
				<i>Scarus frenatus</i>	392	19	1215.9
				<i>Caesio caerulaurea</i>	55	19	1215.9
				<i>Epinephelus polyphemadion</i>	57	21	1216.5
				<i>Chaetodon ulietensis</i>	139	19	1216.8
				<i>Chlorurus sordidus</i>	1049	19	1216.9
				<i>Lutjanus monostigma</i>	84	19	1217
				2 < ΔAIC < 4			
				<i>Aprion virescens</i>	169	20	1217.9

Table S8. Comparative results of potential key species for fish biomass using relative intra-community (left) and intra-specific (right) biomass to define presence (threshold defined as 5%). AIC of the initial model M_0 provides reference Akaike Information Criterion (AIC M_0). Presence of each species has been added to M_0 as binary variable using 2 criteria: i) contribution of species to total biomass higher than 5% and ii) the 95th percentile of the biomass distribution of each species. For each species, number of raw occurrences, number of occurrences meeting the threshold, and AIC of the model accounting for the species (AIC M_1) are reported. For clarity, only key species are presented and those that are significantly related to fish biomass under the 2 approaches are highlighted in grey. Some species did not reach the minimal occurrence required and thus could not be tested using the intra-specific approach (asterisk).

INITIAL MODEL (M0): AIC = 1221.2							
INTRA-COMMUNITY				INTRA-SPECIFIC			
Fish species	Nb of occ.	Nb of occ. (5%)	AIC M1	Fish species	Nb of occ.	Nb of occ. (5%)	AIC M1
<i>Caranx ignobilis</i>	98	81	1169.2	<i>Lutjanus bohar</i>	448	92	1116.8
<i>Lutjanus bohar</i>	448	238	1186.2	<i>Chlorurus microrhinos</i>	314	92	1154.1
<i>Macolor niger</i>	157	48	1207.1	<i>Caranx ignobilis</i>	98	92	1156.2
<i>Platax teira</i> *	22	8	1207.6	<i>Macolor niger</i>	157	92	1164.1
<i>Lethrinus atkinsoni</i> *	74	38	1209.8	<i>Hipposcarus longiceps</i>	163	92	1167.3
<i>Plectropomus laevis</i>	129	86	1210.1	<i>Monotaxis grandoculis</i>	488	92	1171.2
<i>Naso hexacanthus</i>	113	58	1212.7	<i>Naso unicornis</i>	380	92	1183.8
<i>Bolbometopon muricatum</i> *	38	23	1214.5	<i>Acanthurus lineatus</i>	282	92	1186.4
<i>Chlorurus microrhinos</i>	314	167	1215.1	<i>Plectropomus laevis</i>	129	92	1186.9
<i>Lutjanus gibbus</i>	191	72	1215.5	<i>Caranx melampygus</i>	286	92	1191.8
<i>Chaetodon flavirostris</i> *	58	3	1216.4	<i>Ctenochaetus striatus</i>	954	92	1193.9
<i>Elagatis bipinnulata</i> *	32	18	1216.9	<i>Scarus altipinnis</i>	181	97	1194.7
<i>Hipposcarus longiceps</i>	163	41	1217.2	<i>Lutjanus gibbus</i>	191	92	1198.2
				<i>Naso hexacanthus</i>	113	92	1200.1
				<i>Aprion virescens</i>	169	92	1201.1
				<i>Scarus rubroviolaceus</i>	367	95	1204.9
				<i>Cheilinus undulatus</i>	134	92	1205.8
				<i>Pterocaesio tile</i>	181	92	1206.8
				<i>Lutjanus kasmira</i>	202	92	1207.8
				<i>Acanthurus blochii</i>	225	92	1208.5
				<i>Cetoscarus bicolor</i>	144	92	1210.7
				<i>Macolor macularis</i>	107	96	1210.9
				<i>Scarus frenatus</i>	392	92	1211.1
				<i>Zanclus cornutus</i>	532	92	1212.2
				<i>Chlorurus sordidus</i>	1049	92	1212.2
				<i>Heniochus chrysostomus</i>	143	97	1213.6
				<i>Naso lituratus</i>	624	92	1213.8
				<i>Zebrosoma veliferum</i>	295	93	1213.9
				<i>Kyphosus sp.</i>	145	95	1215.2
				<i>Cephalopholis argus</i>	570	92	1215.3
				<i>Acanthurus dussumieri</i>	168	92	1215.3
				<i>Variola louti</i>	139	92	1216.2
				<i>Pomacanthus imperator</i>	137	95	1216.5
				<i>Scarus niger</i>	447	92	1216.9
				<i>Naso brevirostris</i>	155	94	1217.1
				Nb occ. > 90 is required to be tested			
				<i>Platax teira</i> *	22	NA	NA
				<i>Lethrinus atkinsoni</i> *	74	NA	NA
				<i>Bolbometopon muricatum</i> *	38	NA	NA
				<i>Chaetodon flavirostris</i> *	58	NA	NA
				<i>Elagatis bipinnulata</i> *	32	NA	NA

Table S9. Comparative results of potential key species for coral cover using relative intra-community (left) and intra-specific (right) biomass to define presence (threshold defined as 1%). AIC of the initial model M_0 provides reference Akaike Information Criterion (AIC M_0). Presence of each species has been added to M_0 as binary variable using 2 criteria: i) contribution of species to total biomass higher than 1% and ii) the 99th percentile of the biomass distribution of each species. For each species, number of raw occurrences, number of occurrences meeting the threshold, and AIC of the model accounting for the species (AIC M_1) are reported. For clarity, only key species are presented and those that are significantly related to coral cover under the 2 approaches are highlighted in grey. Some species did not reach the minimal occurrence required and thus could not be tested using the intra-specific approach (asterisk), while some species had an AIC lower than M_0 but did not exceed the performance criterion ($\Delta AIC > 4$) to be considered as key species (underlined).

INITIAL MODEL (M_0): AIC = -40.2							
INTRA-COMMUNITY				INTRA-SPECIFIC			
Fish species	Nb of occ.	Nb of occ. (1%)	AIC M_1	Fish species	Nb of occ.	Nb of occ. (1%)	AIC M_1
<i>Acanthurus albipectoralis</i>	24	15	-59.9	<i>Epinephelus fuscoguttatus</i>	11	9	-55.4
<i>Lutjanus bohar</i>	199	172	-57.1	<i>Naso hexacanthus</i>	54	8	-53.1
<i>Bodianus loxozonus</i>	35	7	-53.7	<i>Lutjanus monostigma</i>	35	8	-50.3
<i>Plectorhinchus picus</i>	10	9	-50.7	<i>Lutjanus bohar</i>	199	8	-49.9
<i>Scarus frenatus</i>	158	132	-47.5	<i>Acanthurus albipectoralis</i>	24	8	-47.4
<i>Chlorurus strongylocephalus</i>	32	30	-46.9	<i>Chaetodon reticulatus</i>	86	8	-47.1
<i>Macolor niger</i>	67	50	-46.6	<i>Paracirrhites nissus</i>	24	11	-46.3
<i>Siganus argenteus</i>	20	11	-44.2	<i>Plectorhinchus picus</i>	10	8	-46
				<i>Labroides bicolor</i>	26	14	-45.9
				<i>Thalassoma lutescens</i>	214	8	-44.8
				<i>Hemitaurichthys polylepis</i>	22	8	-44.3
				2 < ΔAIC < 4			
				<i>Chlorurus strongylocephalus</i>	32	8	-35.6
				<i>Bodianus loxozonus</i>	35	6	-36.7
				<i>Scarus frenatus</i>	158	8	-37.3
				<i>Macolor niger</i>	67	8	-37.9
				<i>Siganus argenteus</i>	20	8	-38.1

Table S10. Comparative results of potential key species for coral cover using relative intra-community (left) and intra-specific (right) biomass to define presence (threshold defined as 5%). AIC of the initial model M_0 provides reference Akaike Information Criterion (AIC M_0). Presence of each species has been added to M_0 as binary variable using 2 criteria: i) contribution of species to total biomass higher than 5% and ii) the 95th percentile of the biomass distribution of each species. For each species, number of raw occurrences, number of occurrences meeting the threshold, and AIC of the model accounting for the species (AIC M_1) are reported. For clarity, only key species are presented and those that are significantly related to coral cover under the 2 approaches are highlighted in grey. Some species did not reach the minimum occurrence and thus could not be tested using the intra-specific approach (asterisk).

INITIAL MODEL (M_0): AIC = -40.2							
INTRA-COMMUNITY				INTRA-SPECIFIC			
Fish species	Nb of occ.	Nb of occ. (5%)	AIC M_1	Fish species	Nb of occ.	Nb of occ. (5%)	AIC M_1
<i>Acanthurus albipectoralis</i> *	24	6	-62.7	<i>Chaetodon trifascialis</i>	132	42	-71
<i>Lutjanus bohar</i>	199	131	-55.9	<i>Acanthurus leucosternon</i>	44	38	-63.7
<i>Siganus argenteus</i> *	20	4	-52.1	<i>Chaetodon guttatissimus</i>	40	40	-60.5
<i>Scarus frenatus</i>	158	60	-50.7	<i>Zebrasoma scopas</i>	101	39	-60.4
<i>Macolor niger</i>	67	28	-49.2	<i>Chaetodon baronessa</i>	65	44	-58.6
				<i>Hemigymnus fasciatus</i>	166	38	-58.2
				<i>Chaetodon unimaculatus</i>	48	38	-56.6
				<i>Chaetodon meyeri</i>	40	40	-54
				<i>Chaetodon reticulatus</i>	86	38	-52.8
				<i>Scarus frenatus</i>	158	38	-52.4
				<i>Thalassoma lutescens</i>	214	38	-51
				<i>Macolor niger</i>	67	38	-50.3
				<i>Thalassoma hardwicke</i>	123	39	-48.7
				<i>Lutjanus bohar</i>	199	38	-47.1
				<i>Chaetodon pelewensis</i>	70	63	-46.0
				<i>Naso brevirostris</i>	110	39	-45.8
				<i>Lutjanus gibbus</i>	55	38	-44.7
				<i>Sargocentron caudimaculatum</i>	39	38	-44.3
Nb occ. > 36 is required to be tested							
<i>Acanthurus albipectoralis</i> *	24				24	NA	NA
<i>Siganus argenteus</i> *	20				20	NA	NA

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