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Recovery potential of the world's coral reef fishes 1 2 M. Aaron MacNeil^{1*}, Nicholas A.J. Graham², Joshua E. Cinner², Shaun K. Wilson³, Ivor D. 3 Williams⁴, Joseph Maina^{5,6}, Steven Newman⁷, Alan M. Friedlander⁸, Stacy Jupiter⁶, Nicholas 4 5 V.C. Polunin⁷, and Tim R. McClanahan⁶ 6 7 ¹Australian Institute of Marine Science, PMB 3 Townsville MC, Townsville, QLD 4810 8 Australia 9 ²Australian Research Council Centre of Excellence for Coral Reef Studies, James Cook University, Townsville, QLD 4811 Australia 10 ³Department of Parks and Wildlife, Kensington, Perth WA 6151 Australia 11 ⁴ Coral Reef Ecosystems Division, NOAA Pacific Islands Fisheries Science Center, Honolulu, HI 12 13 96818 USA ⁵Australian Research Council Centre of Excellence for Environmental Decisions (CEED), 14 15 University of Queensland, Brisbane St Lucia QLD 4074 Australia 16 ^oWildlife Conservation Society, Marine Programs, Bronx, NY 10460 USA 17 ⁷School of Marine Science and Technology, Newcastle University, Newcastle upon Tyne NE1 18 7RU UK 19 ⁸Fisheries Ecology Research Lab, Department of Biology, University of Hawaii, Honolulu, HI 20 96822 USA 21 *Correspondence to: a.macneil@aims.gov.au 22 23 24 Ongoing degradation of coral reef ecosystems has generated substantial interest in how management can support reef resilience^{1,2}. Fishing is the primary source of diminished reef 25 function globally³⁻⁵, leading to widespread calls for additional marine reserves to recover 26 fish biomass and restore key ecosystem functions⁶. Yet there are no established baselines for 27 28 determining when these conservation objectives have been met or whether alternative 29 management strategies provide similar ecosystem benefits. Here we establish empirical 30 conservation benchmarks and fish biomass recovery timelines against which coral reefs can 31 be assessed and managed by studying the recovery potential of more than 800 coral reefs 32 along an exploitation gradient. We found that resident reef fish biomass in the absence of 33 fishing (B_{θ}) averages ~1000 kg/ha and that the vast majority (83%) of fished reefs are 34 missing more than half their expected biomass, with severe consequences for key ecosystem 35 functions such as predation. Given protection from fishing in both open and restricted 36 areas, reef fish biomass has the potential to recover within 35 years on average and within 37 59 years when heavily depleted. Importantly, alternative fisheries restrictions are largely 38 (64%) successful at maintaining biomass above 50% of B_0 , sustaining key functions such as 39 herbivory. Our results demonstrate that critical ecosystem functions can be maintained 40 through a range of fisheries restrictions, allowing coral reef managers to develop recovery

plans that meet conservation and livelihood objectives in areas where marine reserves are not socially or politically feasible solutions.

43

44 There is widespread agreement that local and global drivers need to be addressed to reduce the degradation of coral reef ecosystems worldwide^{1,2}. Numerous reef fisheries are so severely 45 overexploited that critical ecosystem functions such as herbivory and predation are at risk $^{3-5}$. 46 Attempts to rebuild reef fish abundances and associated functions require clear timeframes over 47 48 which assemblages can be restored, and viable management alternatives, such as marine reserves 49 or gear restrictions, that promote recovery. Here we develop the first empirical estimate of coral 50 reef fisheries recovery potential, compiling data from 832 coral reefs across 64 localities 51 (countries and territories; Fig. 1a) to: i) estimate a global unfished biomass (B_0) baseline – *i.e.* the 52 expected density of reef fish on unfished reefs (kg/ha); ii) guantify the rate of reef fish biomass 53 recovery in well-enforced marine reserves using space-for-time substitution; iii) characterize the 54 state of reef fish communities within fished and managed areas in terms of depletion against a B_0

baseline; iv) predict the time required to recover biomass and ecosystem functions across the
 localities studied; and v) explore the potential returns in biomass and function using off-reserve
 management throughout the broader reefscape.

58

59 We used a Bayesian approach to jointly estimate B_0 as the recovery asymptote from well-60 enforced marine reserves (where fishing is effectively prohibited; Fig. 1b) and the average 61 standing biomass of unfished remote areas more than 200 km from human settlements (Fig. 1c). 62 We first used a space-for-time analysis of recovery in well-enforced marine reserves that varied 63 in age and controlled for available factors known to influence observed fish biomass, including local net primary productivity, the percentage of hard coral cover, water depth, and reserve size⁶ 64 65 (Fig. 1b). We then modelled B_0 by linking this recovery data with prior information⁴ on B_0 and 66 biomass from remote reefs (Fig 2c), an approach that explicitly assumes that marine reserves 67 have the potential to recover to such levels in the absence of complicating factors, such as poaching or disturbance, and are of appropriate size⁶. Globally, expected B_0 for diurnally active, 68 resident reef fish was 1013 [963, 1469] kg/ha (posterior median [95% highest posterior density 69 70 intervals]), with a biomass growth rate (r_0) of 0.054 [0.01, 0.11] from an estimated initial biomass 71 in heavily fished reefs of 158 [43, 324] kg/ha (Fig. 1). The wide uncertainty in absolute B_0 72 reflected variability in average biomass among remote localities (from ~500 to 4400 kg/ha; log-73 scale coefficient of variation (CV)=0.08; geometric CV=0.61) as well as differences in 74 productivity, hard coral cover, and atoll presence among reefs (Extended Data Fig. 1). We found

no evidence of data provider bias (Extended Data Fig. 2) and model goodness-of-fit was high
 (Bayesian p-value=0.521; Extended Data Fig. 3).

77

The status of reef fish assemblages on fished reefs against a B_0 baseline varied considerably by

79 locality and whether there were management restrictions on fishing activities. Fished reefs (those

80 that lacked management restrictions) spanned a wide range of exploitation states, from heavily

81 degraded in the Caribbean and Western Pacific, to high-biomass in the remote but inhabited

Pitcairn and Easter Islands (Fig. 2a). While previous studies have assessed how global reef fish yields relate to human population density⁷, we characterise, for the first time, the state of fished

- reefs against an empirical baseline. Of concern was that more than a third of the fished reefs
- sampled had biomass below $0.25 B_0$, a point below which multiple negative ecosystem effects of
- overfishing have been shown to occur in the Western Indian Ocean⁷. Only two localities, in
- Papua New Guinea and Guam, were at or near 0.1 B_0 , a fisheries reference point assumed to
- ⁸⁸ indicate collapse⁸. Reef fish assemblages fared far better when fishing activities were restricted in
- some way, including limitations on the species that could be caught, the gears that could be used,
- and controlled access rights (Fig. 2b). None of the localities with fisheries restrictions had
- average biomass levels below $0.25 B_0$ and 65% were above $0.5 B_0$, although some individual
- 92 reefs within localities were below this level (Fig. 2b).
- 93

Despite extensive research into the benefits and planning of marine reserves, there is limited understanding of how long it takes reef fishes to recover once protected from fishing, limiting the

- ability of decision makers to navigate management tradeoffs. To estimate recovery times for
- 97 fished and restricted reefs under hypothetical protection from fishing, we used the empirical
- 98 recovery curve from marine reserves to back-calculate posterior virtual reserve ages (VA_i) for
- 99 each locality, given their estimated level of fish biomass. We estimated the expected age of
- 100 reserves at 90% recovery $(AR_{0.9})$ and subtracted the virtual reserve ages to calculate reef-specific
- 101 expected recovery times $(TR_{0.9,i})$ under full closure (*i.e.* $TR_{0.9,i} = AR_{0.9} VA_i$). By sampling these
- 102 quantities from the posteriors of our Bayesian model, we were able to develop probabilistic time
- 103 frames for management along an expected path to recovery. Consistent with other studies on
- 104 recovery benchmarks⁹, and the United Nations Food and Agricultural Organization's (FAO)
- 105 definition of underexploited fisheries being between 0.8 and $1.0^{(10)}$, we defined recovered at 0.9
- 106 of B_0 , but also estimated median recovery timeframes for a range of other recovery benchmarks
- 107 and rates of increase (Methods).
- 108
- 109 On average, the fished and fishing-restricted reefs surveyed within localities would require 35 110 years of protection from fishing to recover to $0.9 B_0$, while the most depleted reefs would require
- 111 59 years (Fig. 2c; Extended Data Fig. 4). Recovery times depended critically on the estimated
- rate of biomass recovery and the recovery benchmark used (Extended Data Fig. 5). Although the
- influence of marine reserves can be detected within several years¹¹, our global analysis supports
- previous case studies^{12,13} and a meta-analysis¹⁴ showing comprehensive recovery of reef fish
- biomass likely takes decades to achieve. This suggests that the majority of marine reserves,
- having been implemented in the past 10 to 20 years, will require many more years to achieve
- 117 their recovery potential, underscoring the need for continued, effective protection and
- 118 consideration of other viable management options.
- 119
- 120 To understand how the ecosystem functions provided by fishes change with protection from
- 121 fishing, we examined relative changes in functional group biomass along the gradient from
- 122 collapsed (101 [68, 144] kg/ha) to recovered (908 [614, 1293] kg/ha), using generalized additive

123 models to characterise trends. Despite substantial variability in the proportion of each functional 124 group among reefs, clear non-linear trends were present in relative function (Extended Data Fig. 125 6). During initial recovery, functional returns of key low trophic level species increased rapidly, 126 including browsers, scraper/excavators, grazers, and planktivores (Fig. 2d; Extended Data Fig. 7). These are some of the most important ecosystem functions on coral reefs, as browsers and 127 128 scraper/excavators promote coral dominance by controlling algae and clearing reef substrate for coral settlement and growth¹⁵; grazers help to limit the establishment of macroalgae by intense 129 feeding on algal turfs¹⁶; and planktivores capture water-borne nutrients and sequester them to the 130 reef food web¹⁷. Critically, the relative functions of grazers and scrapers/excavators reached 80 to 131 132 100% of their maximum biomass by 0.5 B_0 , while browsers, planktivores, and the three top predator groups (macro-invertivores, pisci-invertivores, and piscivores) increased steadily as 133 134 standing biomass increased toward B_0 . This overall pattern of functional change shows that key 135 herbivore functions can be fulfilled at intermediate biomass levels, rather than solely among

- 136 pristine areas.
- 137

138 Studies across gradients of human population and fishing densities have previously found the

highest absolute losses of herbivores⁵ and predators^{18,19} can occur with relatively low fishing

140 pressure; in contrast, our results show that the greatest functional changes occur when more than

half of total biomass has been removed, supporting previous non-linear relationships between

biomass and function^{4,16}. This disparity likely reflects differences in studying the effects of
 fishing on pristine versus altered reefs - where the apex predators not included in our analysis are

readily removed²⁰ - and differences in socioeconomic conditions that influence reef exploitation

- 145 at specific locations²¹.
- 146

147 Although marine reserves have been widely advocated conservation tools⁴, they can be untenable 148 where people depend heavily on reef-based resources, highlighting the need for management

alternatives to regulate fisheries on reefs. Therefore to complement the use of effective marine

- 150 reserves, we estimated expected biomass given alternative fishing restrictions (Fig. 2e), which
- 151 typically receive less resistance from fishers than marine reserves²². On average, reefs with some
- 152 form of fisheries restriction had biomass 27% higher than reefs open to fishing (Fig. 2a,b).
- 153 Critically, on reefs with bans on specific fishing gears, such as beach seines, or restrictions on the
- 154 types of fish that can be caught, such as herbivores, biomass levels were between 0.3 and 0.4 B_0 ,
- the point at which up to 80% of herbivore function was retained (Fig. 2e). Thus, even simple
- 156 fisheries restrictions can have substantial impacts on fish functional groups that support important
- 157 reef processes. Still greater biomass and functional returns were observed on reefs with access-
- restrictions limiting the number of people allowed to fish a reef, such as family relations, or
- 159 where other forms of established local marine tenure enable exclusion of external fishers²¹.
- 160 Although these management alternatives clearly promote important functional gains relative to
- 161 openly-fished reefs, it is only among well-enforced, long-established marine reserves that
- 162 predation is maximized, more than tripling the function of piscivory present on collapsed reefs.
- 163

164 The continuing degradation of the world's coral reefs underscores the need for tangible solutions that promote recovery and enhance ecosystem functions^{4,23}. Our results demonstrate that well-165 enforced marine reserves can support a full suite of reef fish functions given enough time to 166 167 recover. However, for reefs where marine reserves cannot be implemented, we find that ecosystem functions can be enhanced through various forms of fisheries management. 168 169 Addressing the coral reef crisis ultimately demands long-term, international action on globalscale issues such as ocean warming and acidification²⁴, factors that may diminish recovery 170 171 potential by ~6% over the coming decades (Extended Data Fig. 8). Despite these challenges, a 172 range of fisheries management options are available to support reef resilience and it is likely that some combination of approaches will be necessary for success. Having benchmarks and timelines 173 174 within an explicit biomass context, such as those provided here, increase the chances of agreeing 175 on, and complying with, a mix of management strategies that will achieve conservation 176 objectives while sustaining reef-based livelihoods. 177 178 179 180 References 181 182 ¹Hughes, T.P. *et al.* Rising to the challenge of sustaining coral reef resilience. *Trends Ecol. Evol.* 183 25, 633-642 (2010). 184 ²Graham, N.A.J. *et al.* Managing resilience to reverse phase shifts in coral reefs. *Front. Ecol.* 185 186 Environ. 11, 541-548 (2013). 187 188 ³Dulvy, N.K., Freckleton, R.P. & Polunin, N.V.C. Coral reef cascades and the indirect effects of 189 predator removal by exploitation. Ecol. Lett. 7, 410-416 (2004). 190 ⁴McClanahan, T.R. *et al.* Critical thresholds and tangible targets for ecosystem-based 191 192 management of coral reefs. Proc. Natl. Acad. Sci. U.S.A. 108, 17230-17233 (2011). 193 194 ⁵Bellwood, D.R. *et al.* Human activity selectively impacts the ecosystem roles of parrotfishes on 195 coral reefs. Proc. R. Soc. B 279, 1621–1629 (2012). 196 ⁶Edgar, G.J. *et al.* Global conservation outcomes depend on marine protected areas with five key 197 198 features. Nature 506, 216-220 (2014). 199 200 ¹Newton, K. *et al.* Current and future sustainability of island coral reef fisheries. *Curr. Biol.* 17, 201 655-658 (2007). 202 203 ⁸Worm, B. *et al.* Rebuilding global fisheries. *Science* **325**, 578-585 (2009). 204

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255 Supplementary Information

Is linked to the online version of the paper at www.nature.com/nature.

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265 Author Contributions

M.A.M. conceived of the study with N.A.J.G, T.R.M, S.K.W., and J.E.C; M.A.M. developed and implemented the analysis; M.A.M. led the manuscript with N.A.J.G, J.E.C, and S.K.W. All other authors contributed data and made substantive contributions to the text.

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270 Author Information

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- declare no competing financial interests. Correspondence and request for materials should be
- addressed to M.A.M. (a.macneil@aims.gov.au). This is SERF working group contribution #10.
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- 275
- 276

277 **Figure 1** | Global reef fish biomass among management categories. a. Study (n=832) and 278 prior (n=157) sites, with numbers matching panel c.; **b**. posterior median recovery trajectory 279 (black line) of reef fish biomass among reserve locations (n=45), with 95% uncertainty intervals 280 (UI: grey), 95% prediction intervals (PI: dotted line), estimated initial biomass (white circle with 281 50% (thick line), and 95% (thin line) highest posterior densities), and observed UVC data (green 282 symbols); c. posterior biomass for remote locations (n=22; boxplots; 50% quantiles) with data (grey circles), median B_0 (black line), 95% UI (grey shading), and 95% PI (dotted line) from B_0 283 284 in d.; d. prior (violet), joint informed (dark blue), and uninformed posterior (black line) densities 285 for B_0 .

286

Figure 2 | Coral reef fish responses across the spectrum of potential recovery. Posterior

density proportion of B_0 for **a.** fished (n=23) and **b**. fishing-restricted (n=17) coral reef locations,

shaded from red (collapsed = $0.1 B_0$) to green (recovered = $0.9 B_0$). c. expected times to recovery

290 $(0.9 B_0)$ for fished (circles) and restricted (squares) reefs given full, effective closure. **d.** average

reef fish functional returns from collapsed to recovered; e. median estimated proportion of B_0

among reef fishery management alternatives (black circles) with 50% (thick line) and 95% (thin
 line) uncertainty intervals.

295 Methods

296

297 Reef fish biomass estimates were based on instantaneous visual counts (UVC) from 2096 surveys 298 collected from coral reef slopes (*i.e.* the sloping, windward outer reef, selected specifically to 299 standardize the reef habitat and remove potential bias associated with habitat type) on 832 300 individual reef sites (hereafter 'reef'). All data were collected using standard belt-transects 301 (50*5m or 30*4m) or point-counts (7m radius) between 2002 and 2013, with the bulk of the data 302 (92%) collected since 2006 (Supplementary Table 1). Data from belt transects and point counts have repeatedly been shown to be comparable in estimating fish abundance²⁵ and biomass²⁶. 303 304 Within each survey area, reef associated fishes were identified to species level, abundance 305 counted, and length (TL) estimated to the nearest 5 cm. A single experienced observer collected 306 data for each dataset except the NOAA data from the Pacific where multiple observers operate on 307 every sampling mission. However NOAA has extensive protocols in place to ensure their 308 observers are well trained and follow consistent protocols, ensuring the data are consistent and 309 unbiased. We tested for any bias among data providers (capturing information on both inter-310 observer differences, and census methods) by including each data provider as a random effect in 311 our model (see below), which assumes that there are inherent correlations within datasets that 312 affect the means and associated errors estimated from their data. This analysis showed that there 313 was no bias among data providers and that there is little information present in data provider 314 identities (Extended Data Fig. 2). From these transect-level data we retained counts of diurnallyactive, non-cryptic reef fish that are resident on the reef slope, excluding sharks and semi-315 316 pelagics (Supplementary Table 2). Metadata for the surveys are within the James Cook 317 University research data repository, the Tropical Data Hub (https://eresearch.jcu.edu.au/tdh). 318 319 Total biomass of fishes on each transect was calculated using published length-weight 320 relationships or those available on FishBase (http://fishbase.org). During this process we removed 321 35 transects where divers were mobbed by behaviourally-aggregating species (e.g. Acanthurus 322 *coeruleus*; n=34) or high biomass aggregating species (*i.e. Bolbometopon muricatum*; n=1) that 323 led to potentially unreliable estimates of standing biomass according to the data provider. This 324 truncated dataset was averaged to the reef level (i.e. transects within the same section of 325 continuous reef)²⁷ forming 832 distinct reefs that formed the basic data for our study. The data

were sampled from key coral regions around the world; however, the coral triangle, Brazil, West Africa, and the Red Sea/Arabian Sea regions are not represented. Fish species were assigned to

functional groups based on trophic guilds and dietary information from the literature and

FishBase. A key scale in our analysis was 'locality', defined as reef areas from 10's to 100's of km

that generally correspond to individual nations and map closely onto ranges of human influence²⁷, within which reefs were nested for analysis. In this way our analysis consisted of

three spatial scales: reef, locality, and global. This dataset can be obtained from the senior author

333 upon request (MAM; a.macneil@aims.gov.au) and combined with PyMC code in the

334 Supplementary Methods to replicate our Bayesian hierarchical analysis.

We used the PyMC package²⁸ for the Python programming language to conduct our analysis, 336 running the (Metropolis-Hastings; MH) MCMC sampler for 10⁶ iterations, with a 900,000 337 338 iteration burn in and a thinning rate of 100, leaving 1000 samples in the posterior of each 339 parameter; these long (relative to say, Gibbs sampling) burn-in times are often required with a MH algorithm. Convergence was monitored by examining posterior chains and distributions for 340 stability and by running 5 chains from different starting points and checking for convergence 341 using Gelman-Rubin statistics²⁹ for parameters across multiple chains, all of which were at or 342 343 close to 1, indicating good convergence of parameters across multiple chains.

344

345 We used multiple data sources, including remote areas, asymptotes of well enforced marine

reserves, and prior information, to estimate unfished biomass (B_0) and time for recovery. Remote

areas - defined as having no recent history of fishing and being more than 200 km from human

348 settlement - informed local B_{0l} and global B_0 , given reef-specific covariates x_{nj} thought to

influence standing biomass that were available at the majority of localities. These covariates included local net primary production $(NPP)^{30}$, average proportion of hard coral cover³¹, depth of

included local net primary production $(NPP)^{30}$, average proportion of hard coral cover³¹, depth of survey $(m)^{32}$, and having been collected on an atoll $(0/1 \text{ dummy variable})^{33}$. NPP was calculated

as ensemble mean of estimates based on two NPP algorithms applied on MODIS and SeaWIFS

data (*i.e. Carbon-based Production Model-2* (CbPM2)³⁴ and Vertically Generalized Production

Model (VGPM)³⁵: http://orca.science.oregonstate.edu; mg C / m^2 / day. Each of these reef-

355 specific nuisance parameters were mean centred to offset the reef level observations relative to

- the main focus of our model the B_{0l} estimates.
- 357

To ensure an appropriate sub-model structure was used, we evaluated fits of three potential linear and non-linear relationships (linear, second-order polynomial, and third-order polynomial) for each continuous nuisance parameter. We selected the best-fitting relationship for each nuisance parameter individually based on having the lowest deviance information criteria (DIC) value (Extended Data Table 1) and then compared DIC values of a candidate model set having all combinations of each nuisance parameter to select a final model (Extended Data Table 2). We also examined the posterior residuals for each nuisance parameter sub-model to ensure no

heteroskedascitity was present and that errors were normally-distributed (Extended Data Fig. 9).

366

To recognize potential data provider methodological effects, we incorporated data-provider status in our B_0 estimates by adding a random effect ρ_j for data provider *j* in our Bayesian hierarchical model. These factors were included in a log-Normal hierarchical model for B_0 , given reef-scale observations $y_{il,r}$:

371

	$y_{il,r}$	$\sim N(\mu_{il,r},\sigma_l)$
372	$\mu_{il,r}$	$= B_{0l} + \beta_1 x_{coral,i} + \beta_2 x_{coral,i}^2 + \beta_3 x_{coral,i}^3 + \beta_4 x_{atoll,i} + \beta_5 x_{production,i} + \beta_6 x_{production,i}^2 + \beta_7 x_{production,i}^3 + \rho_j$
	B_{0l}	$\sim N(B_0, \sigma_b),$
373		[1,2,3]
374		

276	1 11 . 6	
376	and weakly-informative priors	2
570	and weaking informative prior.	,

377

$$\begin{array}{l} \beta_{1,...,7} & \sim N(0.0,100) \\ 378 & \sigma_{l,b} & \sim U(0.0,100) \\ \rho_{j} & \sim N(0.0,100). \end{array}$$

$$[4,5,6]$$

379

Because this study built upon previous research conducted in the Western Indian Ocean⁷ we used 380 381 the posterior B_{0} estimate from that study as the prior for our analysis: 382

383
$$B_0 \sim LN(7.08, 0.46)$$
 [7]

384

385 allowing us to build on existing knowledge by directly integrating information between studies. As a check for those averse to building upon previous research in this way, we also ran the full 386 387 model using an uninformative B_{θ} prior, resulting in highly similar inferences, albeit with 388 marginally greater uncertainty than the informed estimates (6.92 [6.52, 7.27] log(kg/ha) 389 informed; 6.82 [6.45, 7.23] log(kg/ha) uninformed), demonstrating that the observed data 390 dominated the prior in our analysis.

391

392 To estimate times to biomass recovery we relied on data from well-enforced, previously fished 393 marine reserves from around the world (Fig. 1a) and used a space-for-time substitution approach, 394 assuming the relationship between reserve age and standing biomass follows a standard logistic 395 regression model and the same reef-scale offset terms as above:

- 396
- 397 398

 $\sim N(\mu_a, \sigma_m)$ $\begin{aligned} y_{i,a} & \sim N(\mu_a,\sigma_m) \\ \mu_a & = \frac{B_0}{1 + (B_0 = \mu_0)/\mu_0} e^{-ra} + \beta_1 x_{coral,i} + \beta_2 x_{coral,i}^2 + \beta_3 x_{coral,i}^3 + \beta_4 x_{atoll,i} + \beta_5 x_{production,i} + \beta_6 x_{production,i}^2 + \beta_7 x_{production,i}^3 + \rho_j \end{aligned}$ 399 [8,9] 400

- 401
- 402

403 Here a is the age of the marine reserve in years; μ_0 is the average initial reserve biomass; and r 404 the average rate of biomass increase. This model is somewhat less hierarchically-explicit than 405 equation [2] due to the scarcity of global marine reserve biomass data, and relies on the key 406 assumption that average reserve potential recovery is consistent, absent the reef-scale effects in 407 the model. Importantly, B_0 is the same as in equation [3] and the linear offsets $\beta_{l_{m,7}}$ the same as 408 in [2], meaning their effects were jointly estimated from both remote and marine reserve data. Therefore B_0 is estimated from both the trajectory of marine reserves through time and from the 409 410 average biomass of all areas defined *a priori* as being remote: B_0 is the asymptote in the reserve component of the model and the global mean in the remote component of the model. μ_0 , the 411 minimum biomass at reserve age zero, was given an uninformative $\sim U(1,10)$ prior that spanned 412

- 413 the range of the data; the standard deviation σ_m was as in [5]; $x_{size,i}$ was set to allow for potential
- 414 effects of reserve size, thought to be an important component of reserve success 6 .
- 415

416 Next we estimated standing reef fish biomass across a range of fished locations, again

417 hierarchically, given observer effects and reef-level observations within each location:

$$\begin{aligned} y_{il,f} &\sim N(\mu_{il,f},\sigma_f) \\ \mu_{il,f} &= B_{l,f} + \beta_1 x_{coral,i} + \beta_2 x_{coral,i}^2 + \beta_3 x_{coral,i}^3 + \beta_4 x_{atoll,i} + \beta_5 x_{production,i} + \beta_6 x_{production,i}^2 + \beta_7 x_{production,i}^3 + \rho_j \\ B_{l,f} &\sim N(0.0,100) \end{aligned}$$

...

- 419
- 420

421 Here the $B_{l,f}$ terms denote independent log-biomass priors per location as we did not assume any 422 parent (hierarchical) structure among locations other than potential data-provider effects; the 423 standard deviation prior for σ_f was as in [5]. Note that fishing pressure is a continuous variable 424 that implicitly underlies the observed differences in exploitation state outside of the factors 425 included in our analysis.

426

427 To estimate the standing biomass across a range of management categories, z, we applied similar 428 methods:

429

430

431

As for the fished locations, the $B_{l,z}$ terms denote independent log-biomass priors per location and the standard deviation prior for σ_z was as in [5]. Management alternative effects were calculated as the average of the location-level posteriors for each group. Note that some locations in the data (Agrihan, Alamagan, Asuncion, Farallon de Pajaros, Guguan, Maug, Pagan, Rose, and Sarigan) were passively fishery-restricted due to isolation limiting effort that could be directed at the resource and, as a trait that cannot be actively managed, we excluded these locations from this section of our analysis.

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- 440

441 Overall model fit442

443 We conducted posterior predictive checks for goodness of fit (GoF) using Bayesian p-values³⁶,

444 whereby fit was assessed by the discrepancy between observed or simulated data and their

expected values. To do this we simulated new data (y_i^{new}) by sampling from the joint posterior of

[10,11,12]

446 447

$$D(\mathbf{y}|\theta) = \sum_{i} (\sqrt{y_i} - \sqrt{\mu_i})^2$$
[16]

or simulated data, given their expected values (μ_i) :

our model (θ) and calculated the Freeman-Tukey measure of discrepancy for the observed (y_i^{obs})

451 yielding two arrays of median discrepancies $D(y^{obs}|\theta)$ and $D(y^{new}|\theta)$ that were then used to 452 calculate a Bayesian p-value for our model by recording the proportion of times $D(y^{obs}|\theta)$ was 453 greater than $D(y^{new}|\theta)$ (Extended Data Fig. 3). For models not showing evidence of being 454 inconsistent with the observed data, $D(y^{obs}|\theta)$ will greater than $D(y^{new}|\theta)$ 50% of the time, giving 455 a p-value = 0.5; for models that showing evidence of being inconsistent with the observed data, 456 $D(y^{obs}|\theta)$ will, by specification, be greater than (or less than) $D(y^{new}|\theta)$ 95% of the time. 457

458

459 *Times to recovery*

460

We capitalized on our integrated Bayesian model to estimate location-specific recovery times for fished and fishery-restricted reefs within the Bayesian MCMC scheme. First we calculated the average reserve age at recovery (*i.e.* 0.9 B_0 : $B_{0.9}$), given the posterior biomass rate of growth *r* and initial biomass of μ_0 (see posterior parameter estimates in Supplementary Table 3):

466
$$AR_{0.9} = \frac{\log\left[\left(\frac{B_0}{B_{0.9}} - 1\right)/\left(\frac{B_0 - \mu_0}{\mu_0}\right)\right]}{-r}.$$
 [17]

467

468 Next we calculated location-specific virtual reserve ages, given their estimated level of log-biomass:

470
$$VA_i = \frac{\log\left[(\frac{B_0}{B_{l,f/z}} - 1)/(\frac{B_0 - \mu_0}{\mu_0})\right]}{-r},$$
 [18]

471

and subtracted this from $AR_{0.9}$ to give an expected time to recovery for each location:

$$\begin{array}{l} 473\\ 474\\ 475 \end{array} = AR_{0.9,i} = AR_{0.9} - VA_i. \tag{19}$$

Because these calculations were conducted within our MCMC scheme they included posterioruncertainties, given the data and our model.

478 479

480 *Variable recovery targets*

- 482 Our choice to define recovery at $0.9B_0$ was based on recent work on recovery in the North Sea⁹
- and being the midpoint at which individual fish stocks are considered underexploited by the

United Nations Food and Agricultural Organization¹⁰. However, to explore how expected time to 484 recovery was dependent on this choice and the estimated rate of biomass growth, we calculated 485 486 average reserve ages at recovery $(AR_{x,v})$ using the median posterior B_0 and μ_0 values (in [17]) while systematically varying the proportion of B_0 defined as recovered (between 0.8 to 1.0) and 487 the rate of biomass growth (between posterior 95% UI range of 0.012 and 0.11). The resulting 488 489 surface plot showed exponential increases in reserve ages at recovery for slower biomass growth 490 rates and higher values of defined recovery due to the asymptotic nature of the logistic growth 491 model used. (Extended Data Fig. 5)

- 492
- 493
- 494 Potential effects of climate change on B_0
- 495

496 A key assumption of the conclusions drawn from our results is that factors affecting total 497 potential B_0 will remain stable through time. Climate projections have been equivocal as to what might happen to tropical fisheries over the coming decades³⁷, primarily due to uncertainty in how 498 production³⁸ and hard coral habitat³⁹ is expected to change, as well as difficulty in modelling 499 500 tropical coastal habitats³⁷. Nonetheless we used the estimated relationships of log-biomass to productivity and hard coral cover (Extended Data Fig. 1) to explore changes in B_0 due to declines 501 502 in both environmental conditions, using the median posterior estimates from our Bayesian 503 hierarchical model. Results showed that by 2040, given an expected 4% loss of primary productivity³⁸ and a 2% annual loss of coral cover³⁹, we would expect to see a 6% drop in B_0 , to 504

- 505 953 kg/ha (Extended Data Fig. 8).
- 506
- 507

508 Log vs. arithmetic scales of estimation

509

510 By adopting a hierarchical approach we, in effect, chose to average over location-specific

differences in order to make global-scale inferences. We elected to model our data on the log-

scale, as per fisheries convention⁴⁰, because it normalized the variance around our hierarchical
 model, greatly improving the precision of model estimates and the convergence of our model fits.

- 514
- 515 A key related point in our analysis is that our posterior calculations for fractions of B_0 were all on

the arithmetic scale, by exponentiating each location-scale estimate and dividing by e^{B0} . To see

- 517 why this makes sense, taking the posterior estimates for log-biomass from Ahus, PNG (4.54) and 518 B_0 (6.92), Ahus would have retained 4.54/6.92 = 0.66 unfished log-biomass but only $e^{4.54}/e^{6.92}$ =
- 519 0.09 absolute biomass. Given that this is the most heavily exploited reef in our database and that

520 fisheries conventions for defining collapsed and recovered are arithmetic, we retained the

521 arithmetic for our posterior calculations.

- 522
- 523

⁵²⁴ Functional returns

525 To understand how relative reef fish function would be expected to vary over the recovery range 526 527 from collapsed (101 [68, 143] kg/ha) through to recovery (908 [614, 1293] kg/ha), we modelled 528 the average biomass of each functional group across this range (*i.e.* log(101) to log(908) kg/ha) 529 relative to their initial biomass values (*i.e.* average biomass of each functional group at log(101) 530 kg/ha). We deemed these relative changes in biomass 'functional returns' because they express 531 relative increases in function that could be expected given log-scale increases in the total biomass 532 of a given functional group on a coral reef. To do this, and allow for expected non-linarites in 533 functional group responses (due to *e.g.* community interactions, resource dynamics etc., the shape 534 of response to which is currently unknown for most functional groups) we fit a series of 535 generalized additive models (GAMs) to the proportion of each functional group over the 536 community recovery range (Extended Data Fig. 6) in models that included the same covariates as 537 our Bayesian hierarchical model (NPP, average proportion of hard coral cover, depth of survey, and having been collected on an atoll). The form of the model was, for each functional group k: 538

539

540

 $y_{il,k} \sim N(\mu_{il,k}, \sigma_k)$ $\mu_{il,k} = \beta_{0l} + f_1(x_{log-biomass,i}) + \beta_1 x_{coral,i} + \beta_2 x_{atoll,i} + \beta_3 x_{production,i}$ $\beta_{0l} \sim N(0.0,100).$ [20,21,22]

541 542

with the smooth function $f_1(x_{log-biomass,i})$ describing the non-linear relationship between 543 observed functional group proportions and total log-biomass. Dividing the fitted GAMs for each 544 545 functional group by the proportion at collapse provided a measure of expected functional return 546 for each group, where a functional return of 2.0 would mean there is twice the log-biomass of a 547 given functional group present compared to initial conditions. The rationale for this approach was 548 that, as our data span the full range from 0.1 to 0.9 B_0 , we did not need to predict outside of the data, but rather uncover the potentially non-linear changes in relative function for each group 549 550 over this range. All GAMs were run using in the GAMM package in R (http://www.r-project.org), 551 using default smooth parameters that provided consistent fits to a per 0.1 log-kg moving average. 552

- 553
- 554

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591 **Extended Data Figure 1** | Nuisance parameter posterior estimates for modelled recovery.

592 Joint Bayesian hierarchical recovery model **a.** prior (flat black line) and posterior (histograms)

- 593 nuisance parameter densities (vertical dotted line at zero) for factors influencing total reef fish
- 594 biomass (kg/ha), including three parameters for a third order polynomial for hard coral cover [*i.e.* 595 Hard coral (1), (2), (3)], an offset for atoll vs. non-atoll, and three parameters for a third order
- 596 polynomial for productivity [*i.e.* Productivity (1), (2), (3)]; **b.** estimated relationship between
- 597 percent hard coral cover and total biomass using posterior median values (blue line), with 99
- 598 samples from the posterior distribution of the parameters in **a**. (thick grey lines) and marginal
- 599 data (black dots; n=832 reefs); c. plot of observed depth and marginal total biomass given the full
- 600 model (no depth effect present): **d.** estimated relationship between atoll (1) vs. non-atoll (0) and
- 601 total biomass, with marginal data (boxplot and black squares); e. plot of reserve size and marginal
- 602 total fish biomass given the full model (no reserve size effect present); f. estimated relationship
- 603 between productivity and total biomass, with marginal data.
- 604

605 **Extended Data Figure 2 | Data provider random effect posteriors.** Bayesian hierarchical

- model posterior estimated effects of data provider identity, including 95% posterior densities 606
- 607 (thin lines), 50% posterior densities (thin lines), and posterior median values (black circles). 608 Results show no apparent bias among data providers, with little information present in provider identities.
- 609
- 610 611 Extended Data Figure 3 | Bayesian p-values for goodness of fit. Discrepancy-based posterior predictive checks for Bayesian hierarchical model goodness of fit. Points represent Freeman-612
- Tukey discrepancy measures between observed and expected values, $D(y^{obs}|\theta)$, and simulated and 613
- expected values, $D(y^{new}|\theta)$. Plot shows high level of agreement between observed and simulated 614
- discrepancies (p=0.521), indicating the model is not inconsistent with the observed data. Labelled 615
- clusters of distinct points reflect various components of the joint model. 616
- 617
- 618 **Extended Data Figure 4** | Posterior expected times to recovery among localities. Bayesian 619 hierarchical model posterior estimated times to recovery $(0.9B_0)$ for fished (green circles) and
- 620 restricted (amber squares) localities around the world. Black lines are 50% highest posterior
- 621 densities and symbols are posterior median values.
- 622

623 Extended Data Figure 5 | Change in expected reserve age at recovery given specified

- 624 recovery target. Change in expected reserve age at recovery (contour lines; in years) given
- specified values for recovery (as a proportion of B_0) and the 95% highest posterior density range 625
- 626 for the rate of biomass growth (r_0) estimated from a joint Bayesian hierarchical model of
- 627 recovery. Expected recovery time from the most degraded locality (Ahus, PNG; posterior
- median: 94 kg/ha) given r_0 (posterior median: 0.054) is 59 years when recovery is defined at 628 $0.9B_0^9$ (blue dot). 629
- 630

631 Extended Data Figure 6 | Average reef fish functional group across a biomass gradient.

- 632 Generalized additive model (GAM) fits to the relative proportion of **a.** excavators/scrapers, **b.**
- browsers, c. grazers, d. detritivores, e. planktivores, f. micro-invertivores, g. macro-invertivores,
- **h.** pisci-invertivores, and **i.** piscivores in community log-biomass for 832 reef slope sites from
- around the world. Grey dots are reef-level observations; blue dots are a 0.1 log-kg interval
- 636 moving average; GAM fits are represented by mean (solid black line) and 95% confidence
- 637 intervals (dashed line) across the full data range. Mean model fits between initial reserve biomass
- 638 and recovered log-biomass (vertical dotted lines) were scaled relative to their values at $0.1B_0$ to
- 639 characterise reef fish functional responses in Fig. 2.
- 640

641 Extended Data Figure 7 | Generalized additive model (GAM) functional returns with

642 uncertainty. Average relative reef fish functional returns in log-biomass across the range from
 643 collapsed to recovered given the GAM fits in Fig. 2d; lines are GAM fits for log-biomass per
 644 collapsed to recovered given the GAM fits in Fig. 2d; lines are GAM fits for log-biomass per

- 644 functional group relative to their average biomasses at marine reserve age zero (estimated initial 645 log-biomass) in Fig. 1; dashed lines are approximate 95% confidence intervals. Data include 832
- 646 individual reefs.
- 647

648 Extended Data Figure 8 | Potential long term changes in B_{θ} under climate change. Response

649 surface (contour lines) for potential change in B_0 (kg/ha) given a plausible range of decline in

average primary productivity (from current 4.7 kg C/ha/day) and coral cover (from current 26%

average hard coral cover). Response surface based on model estimated effects of productivity and hard coral cover on B_{θ} (Extended Data Fig. 1). Current conditions are in the upper right of the

 β_0 (Extended Data Fig. 1). Current conditions are in the upper right of the panel (blue dot); a plausible scenario for 2040 given a 4% loss of primary productivity and a 2%

- annual loss of coral cover would lead to a 6% drop in expected B_0 , down to 953 kg/ha (dot-
- 655 triangle).
- 656

657 Extended Data Figure 9 | Nuisance parameter residual error plots. Joint Bayesian

658 hierarchical recovery model nuisance parameter absolute residuals and residual histograms for **a**.

- 659 percentage of hard coral cover; **b.** having been collected on an atoll; and **c.** average productivity
- 660 in kg C/ha/day. Dashed red lines indicate non-significant linear trends in absolute residuals
- showing no heteroskedasticity was present; blue solid lines show a normal probability
- 662 distribution fit to the residuals, demonstrating appropriate normal sub-model fit.
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672 Extended Data Table 1 | DIC-based model selection for individual nuisance parameter sub-

- 673 models. Joint Bayesian hierarchical recovery model selection for covariates thought to influence
- 674 standing biomass. Candidate models include linear, second-order polynomial (2) and third-order
- 675 polynomial (3) models for each nuisance parameter fit individually (*i.e.* other nuisance
- parameters absent from full model). Models M1-M12 were used to select the best model form for
- 677 each parameter given the lowest DIC value (bold) for each.
- 678 679

680 Extended Data Table 2 | DIC-based model selection for combined nuisance parameter sub-

- 681 models. Joint Bayesian hierarchical recovery model selection for covariates thought to influence
- 682 standing biomass. Candidate models include varying combinations of linear, second-order
- polynomial (2) and third-order polynomial (3) models for each nuisance parameter selected as the
- lowest DIC-valued model in Extended Data Table 1. Models M13-M34 were used to select
- parameters included in the final model given the lowest DIC value (bold). Atoll 1 indicates atoll
- 686 offset was included in the model.



