

Unexpected high vulnerability of functions in wilderness areas: evidence from coral reef fishes

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32 Abstract

High species richness is thought to support the delivery of multiple ecosystems functions and services under changing environments. Yet some species might perform unique functional roles while others are redundant. Thus, the benefits of high species richness in maintaining ecosystem functioning is uncertain if functions have little redundancy, potentially leaving to high vulnerability of functions. We studied the natural propensity of assemblages to be functionally buffered against loss prior to fishing activities, using functional trait combinations, in coral reefs fish assemblages across unfished wilderness areas of the Indo-Pacific: Chagos Archipelago, New Caledonia and French Polynesia. Fish functional diversity in these wilderness areas is highly vulnerable to fishing, explained by species- and abundance-based redundancy packed into a small combination of traits, leaving most other trait combinations (60%) sensitive to fishing, with no redundancy. Functional vulnerability peaks for mobile and sedentary top predators, and large species in general. Functional vulnerability decreases for certain functional entities in New Caledonia, where overall functional redundancy was higher. Uncovering these baseline patterns of functional vulnerability can offer early warning signals of the damaging effects from fishing, and may serve as baselines to guide precautionary and even proactive conservation actions.

65 **1. Introduction**

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Human activities have already induced the collapse of many ecosystems around the world [1– 67 3] and, in combination with climate change, have triggered major reductions in biodiversity 68 globally [3–7]. Beyond the loss of species, there is a growing awareness that the loss of 69 ecological functions may be the most critical consequence of human disturbances on 70 ecosystems [8-11]. This diversity of ecological functions sustains ecosystem services on 71 72 which humanity depends; such as biomass production [10]. Sustaining ecosystem functions 73 requires both high functional diversity, i.e. a large breadth of ecological functions supported 74 by species [12–16], and high functional redundancy, i.e. a large number of species supporting identical functions in the system [17]. 75

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In theory, species richness is thought to maintain a high level of both functional diversity and 77 redundancy, thus ensuring the long term functioning of ecosystems in a fluctuating 78 79 environment [18]. Indeed, high species richness should increase the probability of having both species supporting different functions (functional diversity) and many species supporting the 80 same ecological functions (functional redundancy) [16]. Many experiments confirm this 81 theory, for example demonstrating the vulnerability of ecosystem functioning to species loss 82 [11, 19]. In natural systems, however, the benefits of high species richness to maintaining 83 84 ecosystem functioning have recently been challenged by three patterns. First, at the scale of 85 ocean basins (and using species checklists), some functions exhibit over-redundancy, i.e. are supported by a disproportionately high number of species, while others are realized by few or 86 87 one species only, even in the richest regions [20]. Second, species that support specific or unique traits in ecosystems (no redundancy) tend to be rare due to their low abundance in 88 89 ecosystems [21]. Third, the distribution of species richness and abundance among trophic groups is more critical than simply the number of species to maintain ecosystem functioning 90 91 and services [22]. These patterns demonstrate the importance of preserving both species and abundance within a wide range of functional groups. 92

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Taken together, these results suggest that high levels of species richness and abundance may not insure ecosystems against functional diversity loss as we once hoped, due to the high vulnerability of some functions that lack redundancy. Yet, this hypothesis has never been tested with empirical data in tropical ecosystems with marginal or no exposure to threats, i.e. 98 where species density and abundances should be close to natural baselines. Assessing the 99 vulnerability of ecological functions to threats in such scenarios would reveal the baseline of 100 functional vulnerability, and the extent to which these ecosystems are buffered against even 101 limited local species declines or extinctions.

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103 Protected areas (PAs) are often used to assess ecological baselines against which biodiversity levels in exploited areas are compared [23]. However, recent studies have shown that even 104 105 PAs cannot be considered as true ecological baselines since anthropogenic disturbances typically started long before these areas were established [2, 24-26]. In addition, most of 106 107 these areas are either too small or embedded in areas influenced by human activities and therefore cannot support the full range of ecological functions [23, 27]. As an alternative, 108 wilderness areas, i.e. large areas geographically isolated from humans by natural geographic 109 barriers or with very limited human presence, may provide ecological baselines close to a 110 'natural' status [2, 25, 27]. Indeed, wilderness areas are traditionally viewed as areas featuring 111 exceptional concentrations of biodiversity and abundance [28], albeit potentially suffering 112 from global changes in a near future [29]. 113

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Here, using coral reef ecosystems along a geographic gradient, we propose to test two hypotheses. First disparate wilderness areas tend to host a similar level of functional diversity, uncovering a consistent baseline for the breadth of functions in ecosystems, despite a high turnover in species composition. Second, this common level of functional diversity remains highly vulnerable to species declines or losses due to a disproportional over-redundancy in some functions and a 'natural' lack of redundancy for some critical functions.

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122 Coral reefs are the most diverse marine ecosystems on Earth [30] and support key services for 123 half a billion people, such as food and income [31]. We quantified the baseline vulnerability to fishing of fish functional diversity in coral reef ecosystems across the Indo-Pacific 124 geographic gradient, taking advantage of extensive surveys in French Polynesia, New 125 126 Caledonia and Chagos. These three wilderness areas all benefit from a high level of isolation from humans [32] and a high level of enforcement due to the presence of military forces, 127 thereby limiting illegal fishing activities Since the ecological knowledge to assess the 128 functions carried by individual species is limited, using species functional traits to infer 129

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       functions offers a viable alternative [33]. Here, we assume that species with more diverse
       combinations of functional traits are more likely to support different functions (e.g. [34, 35]).
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          2. Material and methods
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                  (a) Study regions
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      Remote atolls and islands in three regions (Chagos Archipelago, New Caledonia, and French
       Polynesia) were sampled along the Indo-Pacific biogeographic gradient, encompassing 130°
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      of longitude (16,000 km) (ESM, Figure S1). None of these atolls and islands are inhabited:
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      the northern Chagos Archipelago (the Great Chagos Bank, Peros Banhos and Salomon Island)
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       is more than 650 km south of the Maldives and personnel at the Diego Garcia atoll navy base
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      are not permitted to the northern Archipelago; isolated atolls and islands in New Caledonia
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      (Entrecastaux Archipelago, Astrolabe Reef, and Beautemps-Beaupré) are located between 300
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      to 600 km and > 20 hours by boat from the capital Nouméa [25]; two atolls at the southeast
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      end of the Tuamotu Archipelago (Paraoa and Ahunui) are located approximately 950 km from
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      Papeete, the capital of French Polynesia; and the Acteon Group - a cluster of atolls - is located
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      between 200 and 500 km north of Gambier Island, French Polynesia (Figure 1).
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                  (b) Fish surveys
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      Fish data for the Pacific Islands were collected on outer reef slopes using Underwater Visual
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      Census (UVC) along 50 m transects. Briefly, this method involved two divers each recording
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      species identity, abundance and body length [36]. Transects in New Caledonia (18 transects)
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      and French Polynesia (37 transects) were truncated at 10 m wide (10×50m strip transects). In
      the Chagos Archipelago, fishes were surveyed along outer reef slopes using 50 x 5 m strip
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      transects (79 transects). To make surveys comparable among regions, a common area of 500
      m<sup>2</sup> was obtained by randomly aggregating two Chagos transects (250 m<sup>2</sup>). Accumulation
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      curves of species richness were performed for each region to test for potential biases due to
      survey techniques (ESM, Figure S1). Species densities and abundances were estimated for
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       500 m<sup>2</sup> transects and averaged for each region. Sharks and rays were removed from the main
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      species list due to their specific traits, some poaching in Chagos, and difficulties in assessing
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their abundance using UVC [37]. As such, this study focused on 412 fish species belonging to35 teleost families.

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(c) Functional traits and entities

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To estimate functional diversity, we used six functional traits related to major fish attributes:
1) maximum body size, 2) diet, 3) home range, 4) position over the reef, 5) activity, and 6)
gregariousness [20, 38].

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Fish sizes were coded using six ordered categories: 0-7 cm, 7.1-15 cm, 15.1-30 cm, 30.1-50 171 cm, 50.1-80 cm, and >80 cm. Diet was characterized based on the main items consumed by 172 each species, which led to seven trophic categories: herbivorous-detritivorous (i.e., fishes 173 feeding on turf or filamentous algae and/or undefined organic material), macroalgal 174 herbivorous (i.e., fishes eating large fleshy algae and/or seagrass), invertivorous targeting 175 sessile invertebrates (i.e., corals, sponges, ascidians), invertivorous targeting mobile 176 invertebrates (i.e., benthic species such as crustaceans, echinoderms), planktivores (i.e., fishes 177 eating small organisms in the water column), piscivorous (including fishes and cephalopods), 178 and omnivorous (i.e., fishes for which both vegetal and animal material are important in their 179 diet) [20, 39]. Home range was coded using three ordered categories: sedentary (including 180 181 territorial species), mobile within a reef, and mobile between reefs. Position in the water 182 column was coded using three ordered categories: benthic, bentho-pelagic, and pelagic. Activity period was coded using three ordered categories: diurnal, both diurnal and nocturnal, 183 184 and nocturnal. Schooling was coded using five ordered categories: solitary, pairing, or living in small (3–20 individuals), medium (20–50 individuals), or large (>50 individuals) groups. 185 186 This functional traits database was built using information about the ecology of adult life-187 stages available in the literature and according to observations made in the Indo – Pacific by 188 the survey team [38, 40, 41].

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More detailed descriptions linking these traits to ecological processes can be found in the Supplemental Material of published articles [20, 27]. Since all traits were coded using categories, we defined functional entities (FEs) as groups of species sharing the same trait categories. In total, 412 fish species were clustered into 157 different FEs [20, 27].

(d) Fish functional diversity

In order to compare the level of functional richness among the wilderness areas, we measured 197 the functional richness (FRic) of fishes for each region defined as the volume inside the 198 convex hull occupied by species within a functional space [42, 43]. To build a functional 199 space, we calculated pairwise functional distances between species pairs based on the six 200 functional traits using the Gower metric, which allows mixing different types of variables 201 202 while giving them equal weight [44]. A Principal Coordinates Analysis (PCoA) was performed on this distance matrix to build a multidimensional functional space. We retained 203 the first four principal axes (PCs), which faithfully represent the Gower distance between 204 species [42, 43, 45]. 205

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(e) Taxonomic and Functional β-diversity

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In order to determine if wilderness areas host different species and functional compositions across the Indo-Pacific we assessed taxonomic and functional β -diversity among regions. We used the β -diversity partitioning framework based on the Jaccard dissimilarity index [46, 47]. Taxonomic β -diversity (β_{jac}) equals (Eq.1):

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$$\beta_{jac} = \frac{b+c}{a+b+c}$$
 Eq. 1

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where, a is the number of species in both regions A and B, b is the number of species present in region A but not in region B, and c is the number of species present in region B but not in region A.

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To distinguish between the species replacement vs. nestedness components of β -diversity, we decomposed the pairwise Jaccard dissimilarity index (Eq. 1) into two additive components [46, 47]. The replacement component of the Jaccard dissimilarity index (β_{jtu} , Eq. 2) describes species replacement without the influence of richness difference between regions. This index is formulated as follows:

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$$\beta_{jtu} = \frac{2 \min(b,c)}{a+2 \min(b,c)}$$
 Eq. 2

where a, b, and c are the same as in Eq. 1.

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The nestedness component of the Jaccard dissimilarity index (β_{jne} , Eq. 3) is the difference between β_{jac} and β_{jtu} . This index accounts for the fraction of dissimilarity due to richness difference and is formulated as follows:

Eq. 3

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$$\beta_{jne} = \frac{max(b,c) - min(b,c)}{a+b+c} \times \frac{a}{a+2min(b,c)}$$

235

where *a*, *b*, and *c* are the same as in Eq. 1 and 2. The first term in Eq. 3 expresses a measure of richness difference, whereas the second term corresponds to the dissimilarity version of β_{jtu} that is independent of richness difference $(1 - \beta_{jtu}, [47])$.

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240 The functional β -diversity among regions was decomposed into functional turnover and 241 functional nestedness-resultant components following the same framework [48, 49].

Taxonomic and functional β -diversity and their respective components were computed using the R functions from the "betapart" R package (R 2.15.2, R development Core Team, 2012).

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(f) Functional vulnerability to fishing

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Despite the variety of conceptual approaches, there is growing agreement in defining 247 vulnerability as the combination of three components: i) sensitivity, or the susceptibility of a 248 system to threats, ii) exposure, or the level of those threats on a system, and iii) adaptive 249 *capacity*, or the capacity of the system to prepare for and respond to those threats [39, 50, 51]. 250 By analogy, the level of "functional vulnerability" in a given ecosystem relies on i) functional 251 sensitivity, i.e. the extent to which particular traits are more prone to decline in the face of 252 certain threats [52], ii) exposure or the level of threats, and iii) functional redundancy, i.e. the 253 degree to which the same functional traits are supported by many and/or abundant species. 254

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In wilderness areas, exposure to fishing is absent or extremely low [28]. Therefore, the vulnerability to fishing in wilderness areas, termed here as "baseline vulnerability", is solely defined as the combination of species sensitivity to fishing, driven by their biological traits (e.g. size, growth, reproductive capacity), and the level of functional redundancy which is determined by the natural distribution of species density and abundances among functionalentities (ESM, Figure S2).

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The sensitivity of each fish species to fishing was estimated using a fuzzy logic expert system to take into account eight life history characteristics that are linked to species productivity and other factors that make fish species more or less sensitive to fishing [53]. This indicator has accurate predictive capacity [53] and has been widely recognized as a comprehensive and suitable indicator of fish sensitivity to fishing [54]. The sensitivity to fishing was aggregated at the level of FEs by averaging the sensitivity of all species belonging to the given FE. The scale ranged from 0 to 24.

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Functional redundancy is defined as the level of functional equivalence among species in an 271 ecosystem, such that one function may be performed by one or many species, and one species 272 may substitute for another in the latter case [55]. Here redundancy was assessed using three 273 complementary indices. First we used the number of species composing each FE in each 274 region (FR_S) . However, the number of species per FE is only one aspect of functional 275 redundancy. The distribution of species abundances within FEs represents a complementary 276 aspect of redundancy [56-58]. Therefore, secondly we took into account the number of 277 individuals in each FE (FR_{ab}). 278

The mean abundance per 500m^2 of a species *i* in each region was calculated according to the formula:

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$$\overline{ab}_i = \frac{\sum_{i=1}^N n_i}{N}$$
 Eq. 4

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where *N* is the number of transects per region and n_i is the number of individuals of species *i*. Then functional redundancy of a given FE for a given region FR_{ab} was obtained as follows:

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$$FR_{ab} = \sum_{i=1}^{S} \overline{ab}_i$$
 Eq. 5

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with \overline{ab}_i representing the mean number of individuals of species *i* per 500m² and *S* is the number of species in the given FE for that given region.

Thirdly, we computed a redundancy index accounting for both the number of species and their abundances in each FE. We used the Shannon entropy index $FR_{Shannon}$ with the rationale that a given FE will have more redundancy if represented by many abundant species. Conversely, FEs will have low redundancy if represented by few and rare species. Accordingly, in each region, $FR_{Shannon}$ for each FE was computed as:

$$FR_{shannon} = -\sum_{i=1}^{S} p_i \ln p_i$$
 (Eq. 6)

300

with
$$p_i = \frac{\overline{ab_i}}{\sum_{i=1}^{S} \overline{ab_i}}$$
 (Eq. 7)

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where, p_i is the individuals' proportion of species *i* in the FE, and *S* is the number of species per FE.

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We applied the correction derived from the equivalent number of species [59]:

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$$307 \qquad FR_{shannon_EQ} = \exp(FR_{shannon})$$

308

The equivalent number of species is an unbiased measure of Shannon entropy, following Hill's "doubling" property that ensures that the diversity index doubles with the level of diversity, as opposed to non-linear diversity indices that behave counter intuitively [60].

(Eq. 8)

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Quantitatively, the vulnerability of fish FEs to fishing for each region was assessed using a 313 framework based on multi-criteria decision-making (MCDM) and the TOPSIS method 314 (Technique for Order Preference by Similarity to an Ideal Solution). Applied to our specific 315 case, this technique ranks FEs according to their relative distance to the positive and negative 316 ideal solutions, which represent the conditions obtained when the criteria have extreme values 317 [39, 61]. The positive ideal solution (A^+) corresponds to the conditions where sensitivity to 318 threats is minimum while redundancy is maximum (ESM, Figure S3). Conversely, the 319 negative ideal solution (A) corresponds to the conditions where sensitivity to threats is 320 maximum while redundancy is minimum. 321 Functional vulnerability was then expressed as the relative distance to these positive and 322

negative ideal solutions according to Eq. 9:

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$$V_i = \frac{d^+}{d^- + d^+}$$
 Eq. 9

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where V_i is the vulnerability of functional entity *i*, d^+ is the distance to A^+ and d^- the distance to A^- in Euclidean space. The vulnerability index ranges from 0 if the criteria scores correspond to A^+ and to 1 when the criteria scores correspond to A^- [39, 61].

The vulnerability of a FE is high when both the fishing sensitivity of that FE is high and whenredundancy is low [39, 62].

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(g) Mapping functional sensitivity, redundancy and vulnerability in functional space

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The density distribution of functional redundancy, sensitivity and vulnerability within the functional space was estimated using the kernel method with a Gaussian estimation [63]. The smoothing parameter *h* was estimated using the *ad hoc* method, which is the optimum *h* value obtained for the normal distribution [63]:

$$h = \sigma n^{-\frac{1}{6}}$$
 Eq. 10

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342 where, *n* is the number of FEs, and σ^2 being the estimated variance for *x* and *y* coordinates.

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344	$\sigma = 0.5(\sigma_x^2 + \sigma_y^2)$	Eq. 1
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The kernel density estimation was computed using the "adehabitatHR" R package (R 2.15.2,R development Core Team, 2014).

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Functional redundancy, sensitivity and vulnerability were mapped onto the functional space for each region and their match was estimated with the Pearson product-moment correlation.

352 3. Results

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354 (a) Similar level of functional diversity across wilderness areas

356 The greatest number of species and FEs were found in New Caledonia, encompassing 83% of 357 FEs recorded in the three regions (Figure 2). In contrast, French Polynesia showed the lowest number of species (42%) and FE (69%) (Figure 2 and Additional Information in ESM). 358 Overall, 45% of the total FEs (71) were common to the three regions, whereas only 13% (56) 359 of the species were shared (Figure 2 and Additional Information in ESM). 360

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The functional richness in each region (FRic) ranged from 82% of the global functional 362 363 volume (French Polynesia) to 94.4% (New Caledonia) (ESM, Figure S4). The low variability of functional diversity between biogeographic regions is consistent with the weak level of β -364 functional diversity (functions turnover) between regions ranging from 0.14 to 0.18 365 (maximum is 1). Conversely, the β – taxonomic diversity was higher, ranging between 0.66 366 and 0.79 (maximum at 1) (ESM, Figure S5, Table S1 and Additional Information), implying 367 that most functional entities are present independent of species identities. 368

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(b) Heterogeneous distribution of functional redundancy across wilderness areas and functional entities 371

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Despite the overall stability of functional diversity across the biogeographic gradient, the 373 highest functional redundancy of both species per FE (15) and Shannon entropy (17) was 374 found in New Caledonia, whereas Chagos and French Polynesia only reached 8-9 species 375 376 (Figure 3b & Additional Information in ESM). In each region, more than 60% of FEs were represented by only one species, and 12% to 20% of the FEs were represented by only 2 377 378 species (Figure 3a & Additional Information in ESM), showing that the majority of FEs had low species redundancy. 379

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Mapping variation of Shannon entropy across the functional space showed a highly 381 heterogeneous distribution of functional redundancy for each region, with the highest values 382 disproportionally packed into some parts of the functional space (Figure 4a & ESM Figure 383 384 S7), leaving most of the functional space with low or no redundancy. The extreme concentration of functional redundancy in the top left of the figure (Figure 4a) was 385 represented by sedentary small to medium size detritivorous, invertivorous, planktivores and 386 omnivorous feeders (e.g. surgeonfishes, damselfishes, butterflyfishes) (Figure 4a & ESM 387

Figure S6). The top right concentration of redundancy (Figure 4a) was characterized by small
to medium size invertivorous feeders (mobile prey) (see also Additional Information in ESM).

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391 (c) High vulnerability of functional entities in wilderness areas

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Since most of the functional space had low functional redundancy, the variation in vulnerability to fishing was partly driven by sensitivity to fishing (mean Pearson coefficient across regions = 0.93, p<0.001). Variations in functional sensitivity and redundancy across the functional space were independently distributed (mean Pearson coefficient across locations = -0.09, p > 0.1) albeit the most sensitive species systematically showed very low redundancy (ESM Figure S8).

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Medium to large top predators, both solitary and sedentary, such as groupers (top right) and 400 moray eels (top right), as well as mobile and medium sized schooling species such as the 401 Bluefin trevally (Caranx melampygus) and the Great barracuda (Sphyraena barracuda) 402 (bottom right) were highly vulnerable to fishing in the three regions (Figure 4c, 4d & ESM 403 Figure S6) due to their high sensitivity to fishing (Figure 4b) and low redundancy (1 species 404 per FE (Figure 4a). For the same reasons, large invertivores species feeding on mobile prev 405 such as the Humphead wrasse (Cheilinus undulatus), large invertivores (sedentary and 406 mobile), detritivores and planktivores, located in the upper right side of the functional space 407 408 (Figure 4b & ESM Figure S6) showed a high vulnerability to fishing (Figure 4b & Addition Information in ESM). 409

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At the center of the functional space we observed a high fishing vulnerability peak due to the 411 412 presence of two large, mobile, and medium sized schooling species; the bumphead parrotfish (Bolbometopon muricatum), an invertebrate sessile feeder, and the humpback unicornfish 413 414 (Naso brachycentron), a macroalgae feeder (Figure 4b & ESM Figure S6). At the right side of the functional space we observed a second peak of vulnerability to fishing (hence high 415 416 sensitivity coupled with low redundancy) characterized by the bicolor parrotfish (Cetoscarus ocellatus), a large bioeroder [64] and the bluespine unicornfish (Naso unicornis), a 417 macroalgae feeder, both being the only species in their respective FEs and highly sensitive to 418 fishing, explaining their high vulnerability. 419

Conversely, the high diversity of small to medium size herbivorous-detritivorous, 421 422 omnivorous, invertivorous and planktivores species in some FEs (top left of the functional space), contributed to their high functional redundancy, counterbalancing (mean Pearson 423 coefficient across locations of -0.21, p<0.001) their high sensitivity to fishing and making 424 them less vulnerable (Figure 4). The higher functional redundancy in New Caledonia 425 426 decreased the vulnerability of certain FE. For example the functional entity composed of medium size invertebrates feeders (mainly from the triggerfishes family) was composed of 9 427 428 species in NC and 5 in French Polynesia, thereby, in concert with Shannon entropy, reducing its vulnerability 12% compared to French Polynesia. Variation in redundancy and 429 vulnerability across the functional space depicted by the 3rd and 4th axes of PCoA are showed 430 in Figure S9 (ESM). 431

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433 **4. Discussion**

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(a) Homogeneous level of functional diversity and functional vulnerability across the Indo-Pacific biogeographic gradient

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Surprisingly, despite the large differences in species richness and identity among the three 438 regions, we found very consistent patterns of functional diversity along the species richness 439 biogeographic gradient [65], suggesting that the set of ecological functions for the growth and 440 441 persistence of coral reefs are supported independently of species identity and richness, thus serving as a "natural" baseline indicator. As such, the presence/absence of functional entities 442 443 compared to wilderness areas may be a useful indicator to assess ecosystem conditions under human pressure. This result is in line with a previous study on the scale of biogeographic 444 445 regions demonstrating that functions are maintained along a gradient of species richness [20]. Similar consistency was found for functional vulnerability, suggesting that combinations of 446 447 fish traits have quasi-similar levels of redundancy across coral reefs and the biogeographic gradient, albeit New Caledonia had lower levels of vulnerability for several functional entities 448 449 due to higher functional redundancy.

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451 (b) High vulnerability of ecosystem functioning

453 Studies of vulnerability have typically focussed on species rather than ecological functions [39], and often overlook abundance distributions among species [20, 39]. In addition, the 454 assumption that functional diversity in wilderness areas, seen as a reference, is not buffered 455 against extinction due to relatively high functional vulnerability has never been tested. Here 456 457 we filled this gap by assessing the baseline functional vulnerability of fish assemblages to 458 fishing in some of the most isolated coral reef ecosystems across the Indo-Pacific. We found that even in the near-absence of fishing and considering species abundance, most functional 459 460 entities remain highly vulnerable in each wilderness ecosystem since species diversity is overwhelmingly packed into a small set of FEs, leaving most FEs with no redundancy or 461 functional insurance. Importantly, functional sensitivity and functional redundancy are 462 independently distributed among the combinations of traits. As such, redundancy is seldom 463 compensating for the high sensitivity of some traits, the most sensitive ones often having no 464 redundancy. 465

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Our study is complementary to previous macro-ecological investigations showing the 467 heterogeneous distribution of redundancy across FEs [20, 39], and attests that the same 468 pattern translates to the local scale when considering species abundances. This result is not 469 trivial since even FEs with low species redundancy may be preserved if those species are 470 widespread, abundant, and weakly sensitive to threats. This is not the case for fishes on coral 471 472 reefs. Consequently, high biodiversity and abundance of fishes is unlikely to buffer coral reef 473 ecosystems against functional diversity loss. This result is consistent with the observations that marine ecosystem functions and services scale positively and do not saturate with the 474 475 level of species and FE richness within local faunas [66, 67].

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477 Vulnerability can provide early warning signals of fishing pressure, since the processes performed by vulnerable traits are most likely to be the first to decline, even under low 478 479 pressure [68]. Indeed, from a sustainable management perspective this study highlights key vulnerable functional entities that should be under particular conservation measures. For 480 481 example, the bluespine unicornfish (N. unicornis) has been identified has one of the most important macroalgal consumers on coral reefs [69] yet is targetted by commercial, 482 recreational and artisanal fisheries [70]. Macroalgal feeders play a critical role in preventing 483 and reversing coral-algal shifts and the finding of their high baseline vulnerability to fishing is 484 worrying [8, 9]. Despite recent studies demonstrating signs of overexploitation in the Pacific 485

[70], this species is listed as "Least concern" in the Red list of Threatened Species by the International Union for Conservation of Nature (IUCN). This is also the case for top predators such as the Bluefin trevally (*C.melampygus*) and the great barracuda (*S. barracuda*) which are highly sensitive to fishing [71, 72], while the Bluefin trevally has also been found in very low abundance in marine protected areas compared to wilderness [27]. The IUCN Red list criteria could benefit from taking a trait based approach to uncover the species vulnerability from a functional perspective.

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494 **5.** Conclusion

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With biodiversity being lost in the Anthropocene, the need to carefully manage functional 496 diversity has never been greater. The slow decrease in abundance of functionally important 497 species is insidious, despite strong implications for coral ecosystem functioning and resilience 498 [9]. High redundancy can highlight the core processes that may persist even under high levels 499 of fishing pressure since they are highly buffered against local species extinction. While 500 fishing pressure should decrease on the most vulnerable functional entities, it could 501 potentially be directed toward the least vulnerable functional entities highlighted in the 502 functional space. Local adaptive management measures toward those species could focus on 503 gear restrictions, minimum size limits [73], quotas or fishing bans during spawning times [74, 504 75], or modified angling techniques (e.g. [76-78]). However, those species often do not 505 506 represent the same fishing interest due to their low commercial values [79]. Therefore the burden of reducing fishing pressure on the most vulnerable functional entities should be 507 508 shared along market value chains [80], as fisherman targeting certain species operate under incentives determined by trade and the agents involved at both local and international scales 509 510 [81, 82]. Ultimately, a multilayer management approach considering the complexity of the socio-ecological system is necessary to maintain ecosystem functioning [80]. 511

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513 Data accessibility

514 Functional entity fishing sensitivity and vulnerability on Dryad doi:10.5061/dryad.24gs0

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- 516 Author's contributions

- 517 D.M. and L.V conceived the project; S.D., M.K., L.V., L.W., G.M., P.F., N.A.J.G and many
- others collected the data; S.D., D.M. and L.V. prepared the figures; S.D., D.M. and L.V.
- 519 drafted the initial manuscript; all authors contributed to analyses, writing and interpretation.
- 520

521 **Competing interests**

- 522 The authors declare no competing interests
- 523

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781 Figure captions

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783 **Figure 1.**

Sampled coral reef ecosystems across three regions of the Indo-Pacific. Fish were identified
and counted at the outer reefs of remote atolls (blue stars) in Chagos (79 transects), New
Caledonia (18 transects) and French Polynesia (37 transects).

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788 **Figure 2.**

Sharing of species and functional entities between regions. Venn diagram (top) for the
number of species (A) and the number of functional entities (B) for Chagos (orange), New
Caledonia (green), and French Polynesia (blue).

Percentages indicate the proportion of species and functional entities in each region compared
to the total pool. Histograms (bottom) show the proportion (left-axis) and the number of

species (A) or functional entities (B) (right-axis) in each region (color bars), the proportion of $\frac{1}{200}$ unique species (A) or functional entities (B) in each region (black segments on colored bars)

unique species (A) or functional entities (B) in each region (black segments on colored bars),
the number of species (A) or functional entities (B) for each pair of regions, and the number

the number of species (A) or functional entities (B) for each pair of regions, andof species (A) or functional entities (B) common to the three regions.

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799 **Figure 3.**

The levels of functional redundancy in terms of species richness and Shannon entropy across functional entities for the three regions. Distribution of functional entities (in percentage) along a gradient of functional redundancy in terms of the number of species by functional entity in each region (a). Relationships between the number of species per functional entity and the Shannon entropy (expressed as equivalent number of species, Material and Methods) for each functional entity are shown for each region (b). The bottom panel demonstrates the variation of Shannon entropy for a fixed number of species per functional entity.

- 807 #S/FE is the number of species per functional entity.
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809 **Figure 4.**

Mapping redundancy, sensitivity and vulnerability in the functional space for fish faunas of the three regions. The top row (a) shows for each region the distribution of functional redundancy within functional entities, measured using the Shannon entropy, across the functional space. The middle row (b) shows the distribution of sensitivity to fishing across the functional space while the bottom row (c) shows the distribution of vulnerability to fishing in this space, d) position of vulnerable functional entities in the functional space. Colors indicate the level of vulnerability of functional entities and fish shapes were chosen to illustrate the

- main genus of each functional entity (see also ESM, Figure S6).
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