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Demographic plasticity facilitates ecological and economic resilience in a commercially important reef fish — Source link 🗹

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8	Demographic plasticity facilitates ecological and economic resilience in a commercially
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29	Abstract

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1. Variation in life-history characteristics is evident within and across animal populations.

31 Such variation is mediated by environmental gradients and reflects metabolic constraints or

32 trade-offs that enhance reproductive outputs. While generalizations of life-history

33 relationships across species provide a framework for predicting vulnerability to

overexploitation, deciphering patterns of intraspecific variation may also enable recognition

of peculiar features of populations that facilitate ecological resilience.

2. This study combines age-based biological data from geographically disparate populations

37 of bluespine unicornfish (*Naso unicornis*)—the most commercially-valuable reef-associated

38 species in the insular Indo-Pacific—to explore the magnitude and drivers of variation in life

39 span and examine the mechanisms enabling peculiar mortality schedules.

40 3. Longevity and mortality schedules were investigated across eleven locations encompassing

a range of latitudes and exploitation levels. The presence of different growth types was

42 examined using back-calculated growth histories from otoliths. Growth-type dependent

43 mortality (mortality rates associated with particular growth trajectories) was corroborated

44 using population models that incorporated size-dependent competition.

45 4. We found a threefold geographic variation in life span that was strongly linked to

temperature, but not to anthropogenic pressure or ocean productivity. All populations

47 consistently displayed a two-phase mortality schedule, with higher than expected natural

48 mortality rates in earlier stages of post-settlement life. Reconstructed growth histories and

49 population models demonstrated that variable growth types within populations can yield this

50 peculiar biphasic mortality schedule, where fast growers enjoy early reproductive outputs at

51 the expense of greater mortality, and benefits for slow growers derive from extended

reproductive outputs over a greater number of annual cycles. This promotes population

resilience because individuals can take advantage of cycles of environmental change

54 operating at both short and long-term scales.

55 **5.** Our results highlight a prevailing, fundamental misperception when comparing the life 56 histories of long-lived tropical ectotherms: the seemingly incongruent combination of 57 extended life spans with high mortality rates was enabled by coexistence of variable growth 58 types in a population. Thus a demographic profile incorporating contrasting growth and 59 mortality strategies obscures the demographic effects of harvest across space or time in *N*. 60 *unicornis* and possibly other ectotherms with the combination of longevity and asymptotic 61 growth.

62

Key-words: Acanthuridae, coral-reef fish, ectotherm, life span, mortality, unicornfish,
biogeography, growth plasticity, *Naso unicornis*

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67 Introduction

Variation in life-history traits among individuals within a population is prevalent in 68 common, widespread species as it offers flexibility to adapt to a range of environmental 69 conditions (Bradshaw, 1965). In ectotherms, life span often increases with latitude because 70 71 metabolism slows at lower temperatures (i.e., oxygen consumption, protein synthesis and energy flow are reduced; Clarke & Johnston, 2002). This relationship can be explained by the 72 metabolic theory of ecology (Brown, Gillooly, Allen, Savage, & West 2004; Munch & 73 Salinas, 2009), and reflects the longer-term thermodynamic costs of growth (Clarke, 2017). 74 However, variation in traits such as growth performance, reproductive output, and life span 75 76 also occurs within a population due to differences in temperature, resource availability, and 77 competition experienced by individuals. This variation is constrained by trade-offs that 78 optimize lifetime reproductive outputs for individuals and, collectively, for populations 79 (Stearns, 1976; Trip, Clements, Raubenheimer, & Choat, 2014).

80 The growth and condition of individuals within ectothermic populations is largely mediated by the environmental and social conditions to which they are exposed. Fast growth, 81 82 and hence larger body size, generally facilitates a greater competitive advantage and lower predation risk to individuals compared with smaller members of a cohort, thereby reducing 83 84 the overall risk of mortality (Sogard, 1997), but is often associated with reduced life span (Metcalfe & Monaghan, 2003). This somewhat counterintuitive pattern highlights the trade-85 offs that occur among demographic traits (Arendt, 1997). For example, garter snake 86 populations have been shown to support demographic variants that include both fast- and 87 88 slow-growing individuals (Bronikowski & Arnold, 1999; Bronikowski & Vleck, 2010). In these populations, fast-growing individuals had higher rates of mass-specific resting 89 metabolism and mortality, as well as larger body size, earlier maturation, and a high 90 reproductive output. Slow-growing individuals showed the opposite trends of smaller body 91 size, late maturation, and low reproductive output, along with increased longevity. Such 92 variation in traits is also evident in coral reef fishes exposed to marked environmental 93 variation across relatively small spatial scales (10s of km; Gust, Choat, & Ackerman, 2002; 94 Taylor et al., 2018). 95

Under intense natural selection, particular phenotypes will be favoured for survival 96 (Kingsolver & Pfennig, 2007). For traits under disruptive selection, whereby individuals with 97 extreme trait values have the greatest fitness, a wide range of phenotypes will evolve. If the 98 selective forces acting on fast and slow growth stabilize one another (by producing equal 99 lifetime reproductive values, similar to alternative reproductive strategies within 100 populations), then multiple growth types can be retained through high phenotypic variation 101 (e.g., Rios-Cardenas, Bono, & Morris, 2018; Weinstein et al., 2019). Additionally, for long-102 lived species, trade-offs between growth, mortality, and longevity described above (growth-103 104 type dependent mortality) may favour a range of phenotypes when environmental conditions affecting juvenile recruitment and survival vary over time (Warner & Chesson, 1985). 105

Growth-type dependent mortality within populations can result in highly asymptotic 106 length-at-age relationships of populations, with the oldest individuals often having a smaller 107 mean length (slower growth) than younger individuals that are approaching maximum body 108 size more rapidly (Mulligan & Leaman, 1992). This pattern reflects changes in the 109 demographic composition of cohorts across ages and the delayed senescence of slow-growing 110 111 individuals, rather than a true picture of average growth trajectories. Such a bias may also be evident through instantaneous measures of cohort survivorship, which can yield unusual 112 113 patterns of mortality across adult age classes (e.g., Meekan, Ackerman, & Wellington, 2001; Robertson, Ackerman, Choat, Posada, & Pitt, 2005). For such species, an understanding of 114 patterns of growth and mortality within populations is essential for appropriate management, 115 particularly where the species are the target of commercial harvest (e.g., fisheries). 116 Furthermore, such studies also facilitate a greater understanding of processes leading to 117 evolutionary persistence (Mulligan & Leaman, 1992). 118

The bluespine unicornfish *Naso unicornis* is a coral reef-associated species 119 preferentially targeted by fishing across the insular Pacific (Hoey & Bellwood, 2009; Taylor, 120 2019). Several studies and formal stock assessments based on inferred life-history traits have 121 characterized N. unicornis as among the most vulnerable species to overexploitation in 122 multispecies fisheries in the western Pacific (e.g., Houk et al., 2012; Bejarano, Golbuu, 123 Sapolu, & Mumby, 2013; Nadon, Ault, Williams, Smith, & DiNardo, 2015). However, 124 despite the dominance of this species in regional fisheries, there is very limited spatial or 125 temporal evidence of demographic shifts (e.g., decreases in harvested length or proportion of 126 fishery catch) that might result from overfishing (Ford, Bejarano, Marshell, & Mumby, 2016; 127 Houk, Cuetos-Bueno, Tibbatts, & Gutierrez, 2018; Rhodes et al., 2018). Earlier life-history 128 studies of N. unicornis identified an unusual pattern of mortality in both exploited (Taylor, 129

Rhodes, Marshell, & McIlwain, 2014; Andrews et al., 2016) and unexploited populations
(Choat & Axe, 1996) whereby initial adult-stage mortality was high, but then abruptly
decreased, resulting in a two-phase mortality profile across age classes that results in the
atypical combination of high mortality and long life span. The spatial ubiquity of this pattern,
as well as its potential drivers, are unknown.

Here, we compiled a large data set of published and previously unpublished age-based 135 demographic information for N. unicornis from 20 islands and atolls across the Western, 136 Central, and South Pacific and Central Indian Oceans. Our study had two major objectives. 137 138 The first examined the influence of anthropogenic pressure and environmental differences among locations on variability in life span of the species. Prevailing metabolic theory posits 139 that increasing temperature will negatively influence life span (Munch & Salinas, 2009). 140 However, fishery exploitation may have an over-arching effect, whereby any relationship 141 between temperature and life span is obscured by variability in human extraction, given the 142 143 capacity for fishery-induced age truncation in taxa that are long-lived and harvested heavily (e.g., Beamish, McFarlane, & Benson, 2006). The second objective explored patterns of adult 144 145 mortality among locations that ranged across a large gradient of human population densities. We examined the intrinsic biological characteristics (e.g., growth-type dependent mortality) 146 147 that facilitated population-level longevity, variable mortality profiles across age classes, and the evolutionary implications of variable life-history strategies within populations. To 148 achieve this goal, we used an analysis of patterns of otolith increment-width (a proxy for 149 somatic growth) to test the hypothesis that fast growers died young, whereas the slowest-150 growing individuals attained the oldest ages in a population (Mulligan & Leaman, 1992; 151 Metcalfe & Monaghan, 2003). We followed this with a population model that examined the 152 potential contribution of size-dependent mortality towards population-level biphasic mortality 153 profiles. Finally, we evaluated the results in the context of multispecies fisheries management 154 and the ecological significance of variable life-history traits within species. 155

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157 Materials and methods

To assess the generality of life-history characteristics and geographic variation in agebased demography, we collated age-based biological data (i.e., annualized fish ages and associated body length measurements) derived from sectioned otoliths (ear stones) for *N. unicornis* from 20 islands and atolls, geographically summarized into 11 locations across the Indo-Pacific region (Fig. 1). Specimens were collected using fishery-dependent or - independent sampling protocols (Table 1; expanded sampling methods are provided in theSupporting Information).

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166 PREDICTORS OF LIFE SPAN

We estimated the life span N. unicornis at each location using the maximum age 167 derived from sectioned otoliths. Maximum derived age was validated by bomb radiocarbon 168 analysis of micro-milled cores of otoliths of Oahu specimens (Andrews et al., 2016). This 169 method provides a defensible estimate of life span when derived from large sample sizes 170 171 (Kritzer, Davies, & Mapstone, 2001; Hoenig, 2017). Here, we used age data from 2603 individuals from the eleven locations, with a range of 61-533 samples per location (mean 229 172 ± 153 S.D.). On average, this level of sampling yielded 9.2 specimen samples per age class 173 across locations, which falls within the acceptable level of precision (7-10 per age class) 174 proposed by Kritzer, Davies, and Mapstone (2001). We note that sample sizes for four 175 locations (Chagos Archipelago, Mariana Trench Marine National Monument, Northern 176 Mariana Islands, and Yap) fall below this proposed sample level (2.4-4.5 samples per age 177 178 class. However, these four locations are either unfished or had the lowest human density among populated locations (Yap); thus, sampling efforts at these locations, where fishing 179 180 pressure is low to nonexistent, have an inherently higher likelihood of capturing the oldest age classes. Ultimately, this method was used to approximate rather than directly measure life 181 span. Techniques for sectioning and interpreting otoliths not documented in the literature 182 followed Taylor, Rhodes, Marshell, & McIlwain (2014). All otoliths from published studies 183 (Choat & Axe, 1996; Taylor, Rhodes, Marshell, & McIlwain, 2014; Andrews et al., 2016) 184 were re-examined by the lead author for consistency in reading across locations. 185 Measures of environmental variables sea surface temperature (SST; °C) and 186 chlorophyll-a productivity (Chl-a; g C m³) for each location were derived from remotely 187 sensed data. For SST, we calculated the average annual temperature (SST_A), the average 188

temperature from the coldest month of the year (SST_L), and the average temperature from the 189 warmest month of the year (SST_H) using the Global High Resolution Blended SST database 190 from the NOAA ERDDAP website (1-day composite; 0.0879° resolution) from Jan 2006 to 191 May 2011. For Chl-a, we used the Aqua MODIS Global 4 km database to derive the mean 192 Chl-a value across the period Jan 2006 to Jul 2018. Finally, we calculated human density for 193 each location by dividing the 2010 census population by total coral reef area <30 m deep 194 (km²). We then examined the relationships between explanatory variables and life span 195 estimates using linear models, fitted to raw data across locations using least squares 196

estimation. We used single predictor models given the limited number of data points andcompared these models using Akaike's Information Criterion (AIC).

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0 POPULATION DYNAMICS AND DEMOGRAPHIC STRUCTURE

We explored mortality schedules (patterns of instantaneous mortality across age 201 classes) by fitting age-based catch curves to the natural-log-transformed frequencies of age 202 classes for each location (Ricker, 1975). This standard technique linearizes the exponential 203 decay of survivors across age classes, so that the slope of a linear model fitted to the 204 205 descending limb of the curve estimates the instantaneous mortality rate. Age classes with no sampled individuals or those considered not fully recruited to the fishery or scientific 206 sampling (i.e., age classes younger than the peak in the age-frequency distribution) were 207 omitted from the analysis. We interpreted the slope of the linear fit as representing the 208 instantaneous total annual mortality rate (Z). Initial inspection of catch curves revealed non-209 linear mortality schedules at all locations, with residuals conforming to a U-shaped pattern 210 across ages, suggesting a two-phase mortality schedule (Figure S1, Supporting information). 211 212 Consequently, we used a breakpoint analysis (Zeileis, Leisch, Hornik, & Kleiber, 2002) to establish the optimal breakpoint (i.e., age at which mortality schedules shift) for data from 213 214 each location followed by analysis of covariance to test whether the slopes before and after the breakpoint were significantly different. This analysis fitted two independent lines on 215 either side of the optimal breakpoint; therefore, we also modeled mortality using a segmented 216 analysis (Muggeo, 2008) that generates a continuous two-phase model joined at the 217 breakpoint. We note that linear catch curve analysis is based on the assumption of constant 218 annual recruitment. Species of the genus Naso can experience episodes of mass recruitment 219 (e.g., Doherty et al., 2004) which may bias estimates of mortality rates considerably. 220 However, our analysis does not focus on mortality rate values, but rather on mortality profiles 221 222 across the life span of a population. We further expect our results to be buffered by long life spans (ranging 20-50 years among locations) and multiyear collections for seven of the 223 eleven locations (Supporting Information). 224

A two-phase mortality schedule that does not delineate major life phases (e.g., metamorphosis from pelagic phase or onset of sexual maturation) could reflect either a biased sampling protocol or an intrinsic biological phenomenon. Our sampling protocols varied between fishery-dependent (stratified random market sampling) and fishery-independent (spearfishing during research cruises) among populations, and we have no reason to suspect that an aspect of all the sample collections would facilitate a consistent bias. Hence, we set

out to test the hypothesis that initial growth trajectories within populations yielded different 231 life span potentials that resulted in a distinct change in natural mortality rates across age 232 classes. Specifically, we hypothesized that fast-growing fish had shorter life spans than slow-233 growing fish (mechanisms reviewed in Metcalfe & Monaghan, 2003), such that the oldest 234 fish in a population were exclusively derived from slow-growing individuals. Individual 235 growth histories and life spans cannot be determined from length-at-age data alone (Mulligan 236 & Leaman, 1992). Therefore, to test this hypothesis and its feasibility in relation to 237 population dynamics, we (1) examined annual increment width profiles in four of the five 238 239 longest-lived populations (Oahu, Marianas Trench Marine National Monument [MTMNM], Northern Mariana Islands [NMI], and Chagos Archipelago) as a proxy for somatic growth 240 histories of fish (Campana, 1990) and (2) constructed simple population models exploring the 241 effects of variable growth profiles on population dynamics. For the first objective, we 242 selected fish specimens from the smaller (slow grower) and larger (fast grower) ends of the 243 age-specific length distribution, as well as old fish (representatives from the oldest age 244 classes with appropriate otolith clarity for high-resolution measurements) for increment-245 246 width analysis. Slow and fast growers were taken from age classes displaying a high adult growth trajectory, implying that fish could only reach those length-age combinations through 247 248 slower or faster initial growth compared with the rest of the population. Pre-settlement growth (pelagic phase from hatching to reef settlement) and the first ten years of post-249 250 settlement otolith growth were measured and summarized by growth type for each location separately. Our hypothesis posits that slow-growing and old fish would have invariant 251 increment-width profiles across stages in their early life history, whereas increment widths of 252 fast-growing fish would differ across some stage in their early life history. This hypothesis 253 254 stems from studies that show that early growth rate is the best predictor of adult mortality, with slow growth being associated with the longest life expectancy (Metcalfe & Monaghan, 255 2003). Age-specific analysis of variance tests were used to explore differences in increment 256 widths among growth types. 257

For the population model, we interpreted life-history traits based on the upper and lower limits of the growth curve distribution (fast versus slow growers) separately for the Oahu sample and the combined Mariana Archipelago samples (MTMNM and NMI were pooled to provide adequate sample sizes of growth and reproductive output data). Doing so effectively splits the populations into two growth types with associated errors (Table S1, Supporting information), with growth modelled using the von Bertalanffy growth function:

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$$L_t = L_{\infty} [1 - e^{-k(t - t_0)}],$$

eqn1

where L_t is the predicted length at age t (years), L_{∞} represents the asymptotic length, k is the 266 growth coefficient, and t_0 is the hypothetical age at which length is equal to zero. These 267 growth types had considerable overlap in length-at-age profiles, thereby reflecting the 268 continuum of growth rates and the natural variation observed in the populations on which our 269 models were based. Life-history theory suggests that variable growth types can be maintained 270 within a population if reproductive output is approximately the same among the growth types. 271 272 Our model assumed that only slow-growers reached the oldest age classes and we chose a maximum age for fast growers informed from the breakpoint analysis from mortality profiles 273 274 and overall life span (average breakpoint at $\sim 40\%$ of lifespan). Natural mortality rates (M) for both growth types were estimated based on established relationships with life span (Then, 275 Hoenig, Hall, & Hewitt, 2014; M = 4.899 [maximum life span]^{-0.916}), presuming maximum 276 ages of 50 years (Oahu, slow growers), 20 years (Oahu, fast growers), 40 years (Mariana 277 Archipelago, slow growers), and 15 years (Mariana Archipelago, fast growers). Population 278 decay across age classes was calculated using the standard exponential decay equation: 279 $N_{t+1} = N_t e^{-M}$ 280 281 ean2

where *N* represents population size, *t* represents age class, and *M* represents the instantaneous natural mortality rate. To estimate reproductive output, we then modelled the relationship between fork length and female ovary weight (*OW*; Supporting Information) and multiplied the length-specific *OW* by cohort survivorship (the relationship between *OW* and fork length did not differ between growth types based on empirical data). The ovary weight × survivorship values were summed across age classes from the age at maturity (t_{mat}) to the maximum age (t_{max}) and considered a proxy for total reproductive output (*RO*) of a cohort:

290

$$RO = \sum_{t = t_{mat}}^{t_{max}} N_t O W_t$$

eqn3

In this scenario, we therefore used total mass of propagules through a cohort's existence to
represent reproductive output. We acknowledge that changes in egg size or spawning
frequency across different body sizes may affect actual reproductive value, but this
information was not available for any population in this study. The model assumed an equal
number of fast- and slow-growing recruits in a cohort. Finally, we included an additional

mortality factor (percentage relative to *M*) for slow-growing recruits during the age classes
that overlapped with the life span of fast growers, to represent increased mortality from
competition with faster-growing individuals and increased predation risk. We used an
optimization procedure on this additional mortality factor to find the scenario that yielded
equal lifetime reproductive output estimates from fast-growing and slow-growing
components of a cohort across their respective life spans.

- 302
- 303 **Results**

304 We found a remarkably strong relationship between SST and life span, with the average temperature of the coldest month (SST_L) being the predictor that explained the most 305 variation (Fig. 2). Although life span estimates ranged widely from 16 to 52 years, SST_L 306 explained 83% of the variation across locations ($F_{1,9} = 44.82$, P < 0.0001, AIC = 68.52). All 307 locations were oceanic island systems except for Lizard Island, Great Barrier Reef, which 308 309 was a high island on a continental shelf. Sampling at Lizard Island provided an outlier in the data set, and removal of this point improved the correlation to 93% of variance explained 310 $(F_{1,8} = 111.34, P \le 0.0001, AIC = 54.56)$. Human density (a proxy for anthropogenic 311 pressure) had no relationship with life span ($F_{1,9} = 3.88$, P = 0.0804, AIC = 84.26). Chl-a 312 (natural log-transformed) also did not correlate with life span ($F_{1,9} = 0.44$, P = 0.5234, AIC = 313 87.67). 314

Despite the variability in life span, we observed two-phase mortality schedules of 315 varying intensity in all populations, whereby a breakpoint occurred somewhere between 7 to 316 20 years (Fig. 3). The change in slope before and after the optimal breakpoint was significant 317 at all of the locations (Table S2, Supporting information), with the exception of the Northern 318 Mariana Islands. The instantaneous mortality rate prior to the breakpoint ranged from 1.5 to 319 24.8 times higher than the later part of the life span across locations (comparing disjunct fits 320 here as they are not constrained by one another). The difference in slopes between parts of 321 the mortality schedule was not significantly related to temperature (linear model; $F_{1,10} = 2.05$, 322 P = 0.19), anthropogenic pressure ($F_{1,10} = 0.06$, P = 0.82), or Chl-a ($F_{1,10} = 0.45$, P = 0.52). 323

Increment-widths (Fig. 4a) were derived for slow-growing, fast-growing, and old fish (Fig. 4b) across the first ten years of life. For the four populations examined, increment-width profiles were very similar during pre-settlement growth and across the first ten years for slow-growing and old fish, but growth during year 1 was significantly greater for fastgrowing fish (Fig. 4c). Although fast-growing fish also had significantly larger increment widths for other age classes at Oahu and the Northern Mariana Islands, the large and consistent disparity at year 1 across populations confirms that fast-growing fish attained the
majority of their growth advantage in the first year after settlement (i.e., growth potential is
not predetermined during the pelagic phase). Further, the comparisons of increment-width
profiles between growth types and old fish provide strong support for the hypothesis that the
oldest fish are represented by survivors among the slowest-growing individuals, thus that
fast-growing fish die young.

Our population models independently integrated two growth extremes based on 336 empirical data from Oahu and the Mariana Archipelago. Fig. 5a displays the simulated 337 338 length-at-age information for each growth type for the Oahu population. By integrating the relationship between ovary weight and fork length (Fig. 5b) across natural population decay, 339 we determined that fast-growing and slow-growing cohorts can reach the same lifetime 340 reproductive value when intraspecific competition and predation increase natural mortality of 341 the slow growers by approximately 60% at Oahu and 35% in the Mariana Archipelago. This 342 scenario results in a two-phase mortality schedule similar to the low end observed from 343 empirical data, whereby natural mortality of all growth types combined is nearly doubled 344 345 during the life span of the fast-growing fish (Fig, 5a; Fig. S2, Supporting information). Because fast-growing fish mature earlier and reach larger sizes faster, their reproductive 346 347 output is much greater than slow-growing fish in the first ten years (Fig. 5c; Fig. S2, Supporting information). 348

349

350 4 | Discussion

Contemporary tropical fisheries assessments are often data-poor and based on 351 assumed geographically invariable values for life-history traits. Here, we showed two 352 phenomena that influenced the life history of the commercially-important bluespine 353 unicornfish, N. unicornis, that challenge such assumptions. We found a significant and strong 354 relationship between sea surface temperature and life span, whereby warmer water yielded 355 shorter-lived populations. This is predicted by metabolic theory and supports previous 356 evidence demonstrating high variability in longevity strongly linked to ambient temperature 357 in other acanthurid reef fishes and more generally in ectotherms (Robertson, Ackerman, 358 Choat, Posada, & Pitt 2005, Munch & Salinas, 2009). Failing to account for this relationship 359 in fishery models results in gross over- or under-estimates of exploitation, mortality, and 360 yield. The second phenomenon was the presence of a two-phase mortality schedule, common 361 across all locations examined. This suggests that metabolic trade-offs between growth rates 362 and life spans are conserved within populations irrespective of geographic location. 363

The wide variation in life span recorded across a latitudinal gradient is consistent with 364 the metabolic theory of ecology (Brown, Gillooly, Allen, Savage, & West, 2004; Munch & 365 Salinas, 2009), and is reflective of trade-offs between thermally-enhanced growth and 366 reproductive life span (Trip, Clements, Raubenheimer, & Choat, 2014). Other studies have 367 found a similar threefold change in life span of primary consumers that is tightly correlated 368 with temperature (Robertson, Ackerman, Choat, Posada, & Pitt, 2005; Taylor, Trip, & Choat, 369 2018). For the ocean surgeonfish Acanthurus bahianus, temperature had a much stronger 370 influence on life span than it did on either growth rate or body size, but all three were 371 372 inversely related to temperature (Robertson, Ackerman, Choat, Posada, & Pitt, 2005). These authors also highlighted trade-offs with reproduction, whereby frequency of spawning 373 activity generally decreased with declining temperature. Despite the broad geographic 374 patterns documenting decreased life span with increased environmental temperature in both 375 ectotherms and endotherms, the proximal mechanism underlying variation in life span is still 376 debated (Speakman, 2005). Analysis of ectothermic species, however, consistently indicates a 377 complex integration of thermal constraints on growth in which higher temperatures at low 378 379 latitudes elicit faster growth through more efficient protein synthesis (Clarke, 2017) despite 380 the physiological costs of higher metabolic demands on life span (Metcalfe & Monaghan, 381 2003).

Our results support the hypothesis that intraspecific variation in growth rates within 382 cohorts of *N. unicornis* drives observed patterns of biphasic mortality. Differential growth 383 rates in the pre-maturational and early reproductive phases of the benthic life cycle – as 384 observed through otolith increment width patterns – were clearly linked to variation in 385 mortality rates and maximum life spans in the sampled populations. Individuals with slower 386 initial growth rates lived longer than those with faster initial growth rates. Differential growth 387 rates within a cohort will have impacts on reproductive demography, as fast growth is 388 389 associated with earlier maturation and greater reproductive outputs compared with slow growth in a cohort. Moreover, enhanced growth rate and the associated increases in size 390 would confer advantages in terms of intra-specific competition and reduced predation risk 391 compared with other members of a cohort. However, fast growth and the resulting large body 392 size require greater metabolic demands (i.e., energy expenditure), and although these features 393 generally yield greater reproductive output and a competitive advantage, greater metabolic 394 demands eventually bring a growth cost resulting in early senescence (Speakman, 2005). 395 Ultimately, when selection pressures yield benefits for different growth types, a wide variety 396

of phenotypes can be retained within populations. This concept in population demography isanalogous to portfolio effects in ecological systems.

Our population model implies that a species will only evolve biphasic mortality 399 schedules if slow-growing fish are subjected to a factor that induces additional mortality 400 during the period when they co-occur with shorter-lived fast growers. This additional factor is 401 on the order of a 30-60% greater mortality rate (relative to natural mortality predicted from 402 life span) and presumably stems from lower capacity for successful competition and greater 403 predation risk among smaller-bodied individuals. Empirical field tests support this range as a 404 405 realistic estimate. For instance, demography of newly settled fairy basslets (Gramma loreto) on patch reefs was heavily influenced by the density of older/larger individuals (Webster, 406 2004). Greater abundance of larger conspecifics both reduced feeding rates (by up to 60%) 407 and increased predation risk of juveniles, with mortality doubling in the presence of high 408 adult densities (Webster & Hixon, 2000; Webster, 2004). Hixon & Jones (2005), in a multi-409 410 year study, found that intraspecific competition was the primary driver of density-dependent mortality. Within a cohort, density-dependent effects were lagged across time, only emerging 411 412 once body-size distributions widened because of variability in individual growth profiles, suggesting size-based competitive advantages. Overall, patterns of mortality dependent on 413 414 body size appear to be common among marine fishes, with extreme examples demonstrating greater survival of larger-bodied conspecifics by factors up to nearly 40 (Sogard, 1997). 415 The demographic benefits for fast-growing individuals of earlier maturation and 416

greater competitive abilities within the cohort are properties that might be beneficial if 417 sequential cohorts are confronted by a habitat characterised by a high degree of patchiness on 418 a local scale. Under these circumstances, individuals could exploit patches of high quality 419 food resources, with rapid growth and early maturation leading to enhanced reproductive 420 outputs. Empirical measurements of larval dispersal show high levels of larval retention 421 422 regardless of a species' early life-history characteristics (Jones, 2015). Such retention would ensure that successful offspring would be able to take advantage of the conditions that 423 424 provided for increased reproductive output in the parental generation, whereas the long larval pelagic duration of *N. unicornis* (~71-75 d; Doherty et al., 2004; Andrews et al. 2016) affords 425 the potential to exploit new and distant habitats (Horne, van Herwerden, Abellana, & 426 McIlwain, 2013). However, the oceanic and near reef conditions that strongly influence the 427 success of larval cohorts may vary considerably on decadal scales, and are implicated in 428 generating highly successful year classes (Bergenius, McCormick, Meekan, & Robertson, 429 2005). This is supported in reef fishes of the genus *Naso* that are characterised by episodes of 430

mass recruitment (Doherty et al., 2004). Longevity at the scale observed in acanthurid fishes 431 (3-4 decades post maturation) provides the opportunity for some offspring to experience the 432 oceanic conditions that are outstandingly favourable to larval success that occur at decadal 433 time scales. This represents a variant of the storage effect (Warner & Chesson, 1985) in 434 which slow-growing but long-lived individuals may secure the benefits of favourable 435 conditions for larval settlement and recruitment that occur only episodically. The argument 436 predicts a trade-off between rapid initial growth and maturation versus slower growth with an 437 extended life span that provides opportunity to contribute to numerous annual spawning 438 439 episodes (Metcalfe & Monaghan, 2003), with both variants retained within discrete geographic populations. 440

It remains unclear if the proximal driver of growth variation among N. unicornis 441 populations is inherently genetic, reflects the environmental variability to which individuals 442 are exposed, or represents both factors to some spatially or temporally variable extent. At 443 present the information that would allow us to determine if the different demographic modes 444 represent historical divergences due to adaptation are elusive. However, the increasing 445 446 evidence of fine-scale demographic variation in other fishes and the possibility that this may reflect ecological speciation (Roney, Oomen, Knutsen, Olsen, & Hutchings, 2018) shows that 447 448 the mechanisms underlying intra-population diversification of life histories in reef fishes deserve further investigation. 449

Life-history traits have proven to be powerful predictors of vulnerability to 450 overexploitation in a wide variety of harvested fishes (Jennings, Greenstreet, & Reynolds, 451 1999; Abesamis, Green, Russ, & Jadloc, 2014; Taylor, Houk, Russ, & Choat, 2014). This 452 pattern reflects the highly correlated relationships between various traits across closely-453 related species (Roff, 1984) and is based largely on fundamental principles concerning 454 population replenishment and turnover time. However, initial age-based summaries of 455 surgeonfishes demonstrate a surprising lack of consistent patterns among demographic traits 456 across species (Choat & Robertson, 2002). Surgeonfishes of all genera are long-lived (>30 457 years) and most display 'square' growth profiles (long life span with rapid initial growth that 458 quickly reaches an asymptotic size; Trip, Craig, Green, & Choat, 2014). Given this pattern 459 460 and the relative paucity of empirical derivations of mortality for surgeonfishes, we suspect that the two-phase mortality schedule and demographic characteristics yielding it are not 461 unique to *N. unicornis*, but are presently unappreciated and much more general across 462 acanthurids and possibly other lineages displaying 'square' growth trajectories and extensive 463 longevity. We further postulate that this characteristic facilitates greater species-specific 464

resilience to fishery exploitation than previously presumed, given that empirical derivations
of natural mortality based on observations of maximum life span will overestimate population
turnover time. Hence, while life-history traits are often useful predictors of vulnerability to
overexploitation, we predict that interspecific relationships in traits will ultimately be poor
predictors of yield in fisheries for the acanthurids and other demographically similar species.

Although demographic resilience is certainly a positive feature, we highlight that 470 growing human populations, increases in fishing efficiency resulting from constant 471 advancements in technology, and documented declines in other harvested taxa are causing 472 473 unicornfishes and other surgeonfishes to be harvested at greater rates than ever before (Rhodes et al., 2018). This justifies legitimate concerns regarding both regional and broad-474 scale sustainability. Based on contemporary data sets, the bluespine unicornfish appears to 475 remain a large component of fishery harvests rather than diminishing over time from high 476 fishing pressure, but that does not mean populations are not impacted in other ways. More 477 elusive concerns are the 'Darwinian' effects of sustained fishing pressure (Conover, 2000), 478 whereby strong selection pressure on particular traits (for example, overrepresentation of fast 479 480 growers in fishery harvests) by human harvesting systematically removes genotypes from the gene pool or might impair recovery by indirectly inflating rates of natural mortality on the 481 482 smaller survivors of fishery extraction (Meekan, McCormick, Simpson, Chivers, & Ferrari, 2018). As this effect would require many generations to materialize and consistent surveying 483 to identify, we are at present generally unable to observe the direct or indirect consequences 484 of fishery-induced evolution in the context of coral reef fisheries or ecosystems (Kuparinen & 485 Merilä, 2007). 486

Non-extractive anthropogenic impacts such as warming water temperatures resulting 487 from climate change loom in the future for species like bluespine unicornfish whose life 488 histories are strongly influenced by temperature. Numerous studies have demonstrated the 489 490 diverse physiological, ecological, and interactive responses of fishes and other ectotherms to increases in water temperature within the scope of global warming (Munday, Warner, Monro, 491 Pandolfi, & Marshall, 2013; Morrongiello, Sweetman, & Thresher, 2019). Body sizes will 492 generally decrease due to ocean warming and reduced oxygen levels (Cheung et al., 2013), 493 494 but one cannot predict the effect of generally higher temperatures and lower oxygen on biphasic growth, for example, across populations of this or any other species. Changes in 495 vital rates associated with body size, in addition to those resulting from increased human 496 extraction, will however almost certainly occur in the oceans of the future. 497

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512

513 AUTHORS' CONTRIBUTIONS

B.M.T. and J.H.C. conceived the idea for the study; B.M.T., J.H.C., E.E.D., A.S.H., K.L.R.,
and M.G.M. secured funding for field collections; all authors collected field data; B.M.T.
conducted laboratory and data analysis; B.M.T. led the writing of the manuscript and all
authors contributed critically to drafts and gave final approval for publication.

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519 DATA ACCESSIBILITY

520 Data available in the Dryad Digital Repository: https://doi.org/10.5061/dryad.v2g567n
521 (Taylor et al., 2019).

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721	Table 1. Summary of demographic samples for	or Naso unicornis and associated explanatory factors from eleven locations across the Indo-Pacific

722 region.

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Location	n†	FL range	Age range	Collection method	Winter SST	Productivity (g C m ³)	Human density	Previous references
		(mm)	(years)			(g)	(ppl/km²	
							reef)	
Chagos Archipelago	61	235 - 544	2 - 26	Fishery-independent, research cruise collections	27.792	0.111	0	-
Mariana Trench Marine	96	165 - 520	1 - 37	Fishery-independent, research cruise	25.855	0.053	0	-
National Monument‡				collections				
Northern Mariana Islands§	141	192 - 524	1 - 33	Fishery-independent, research cruise collections	26.905	0.051	0	-
Saipan, CNMI	447	108 - 532	1 - 21	Fishery-dependent, market biosampling	27.478	0.055	524	Sundberg et al. (2015) [method]
Tinian, CNMI	144	222 - 488	2 - 22	Fishery-dependent, market biosampling	27.541	0.055	165	Sundberg et al. (2015) [method]
Guam	248	114 - 520	0 - 23	Fishery-dependent, market biosampling	27.722	0.048	1163	Taylor, Rhodes, Marshel and McIlwain (2014)
Yap, FSM	76	136 - 514	1 - 17	Fishery-independent, research sampling	28.313	0.059	70	-
Lizard Island, Great Barrier	209	60 - 542	0 - 30	Fishery-independent, research sampling	24.876	0.288	0	Choat and Axe (1996),
Reef								Choat and Robertson (2002)
Pohnpei, FSM	232	207 - 496	1 - 16	Fishery-dependent, market biosampling	28.640	0.064	230	Taylor, Rhodes, Marshel and McIlwain (2014)

	Oahu, Hawaii	533	67 - 596	0 - 52	Fishery-dependent, market biosampling	23.726	0.081	2923	Andrews et al. (2016),
									DeMartini, Langston, and
									Eble (2014)
	Tutuila, American Samoa	328	124 - 535	1 - 19	Fishery-dependent, market biosampling	27.930	0.050	1118	Sundberg et al. (2015)
									[method]
723									

- 723
- FL, fork length; SST, sea surface temperature; FSM, Federated States of Micronesia; CNMI, Commonwealth of the Northern Mariana Islands
- 725 *†*Sample size of individuals with age estimates
- 726 ‡Islands of Uracas, Maug, and Asuncion
- 727 §Islands of Alamagan, Pagan, Guguan, Sarigan, Anatahan

Author **N**

728 Figure Legends

Figure 1. Indian and Pacific Ocean study sites where demographic samples of bluespine

- violational violation view of the second sec
- 731 Monument (comprises islands of Farallon de Pajaros [Uracas], Maug, and Asuncion); NMI =
- 732 Northern Mariana Islands (comprises sampled islands of Pagan, Alamagan, Guguan, Sarigan,
- and Anatahan).

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Figure 2. Relationship between maximum derived age (life span) of *Naso unicornis* and mean winter sea surface temperature (SST_L). Temperature was calculated as the mean SST of the coldest month for each location, based on remotely-sensed satellite data from 2006-2011. The relationship ($R^2 = 0.83$) is described by *Maximum derived age* = -6.30*(*SST_L*) + 197.01 and is displayed with 95% confidence bands.

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Figure 3. Age-based catch curves for *Naso unicornis* from all study locations, demonstrating
a ubiquitous two-phase mortality schedule punctuated by a breakpoint. White circles
represent estimates for age classes not fully recruited to the sampling strategy, black circles
represent estimates across the first stage of the mortality schedule, grey circles represent
those for the second stage of the mortality schedule. Optimal breakpoints were computed
using a structural change function. Black lines are continuous segmented fits, grey lines are
disjunct fits.

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Figure 4. (A) Annual increment widths follow measurements along transverse sections of 749 otoliths in Naso unicornis from Oahu, Hawaii (with inset showing the annotated transverse 750 751 section of a fish estimated as 9-yr old), two locations from the Mariana Archipelago (inset map), and the Chagos Archipelago. (B) Increment widths were measured from subsampled 752 'fast-growing', 'slow-growing', and 'old' fish based on length-at-age growth plots. (C) 753 Increment width profiles showing divergence among growth types and 'old' fish across the 754 first ten age classes. 0 = from hatching to reef settlement (pelagic phase), 1 = first year of 755 growth (from settlement to first ring deposition), 2 = second year of growth, 3 = third year of 756 growth, etc. *represents significant differences between growth types (F = fast growers, S =757 slow growers, O = old fish). 758

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Figure 5. (A) Simulated growth profiles (with associated von Bertalanffy growth curves) and
life spans of 'fast-growing' and 'slow-growing' *Naso unicornis* based on the Oahu sample.

Bold lines represent the simulated two-phase mortality schedule (natural mortality rate [*M*] is nearly double in the first phase versus the second phase) from this population and relates to the y-axis. (B) Empirical relationship between ovary weight and fork length for female *N*. *unicornis* from Oahu. (C) Relative and cumulative reproductive value of 'fast-growing' and 'slow-growing' cohorts over their respective life spans.

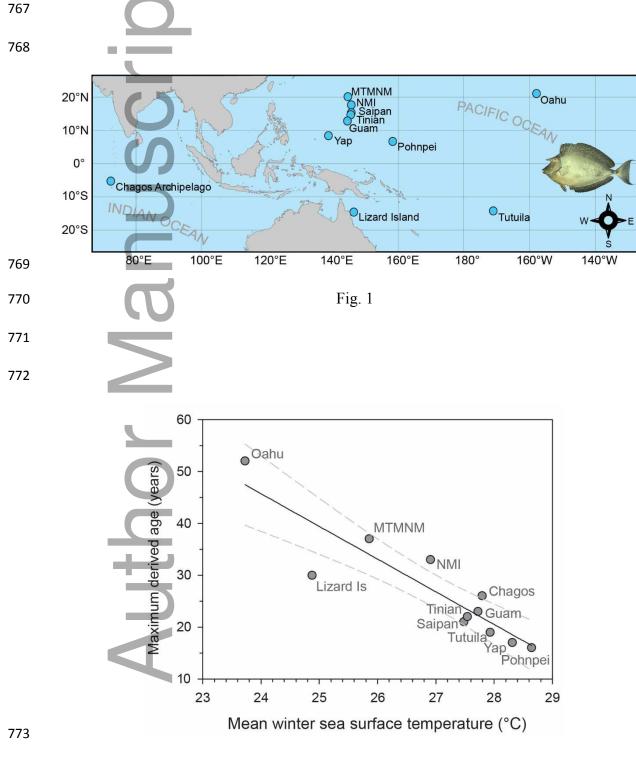
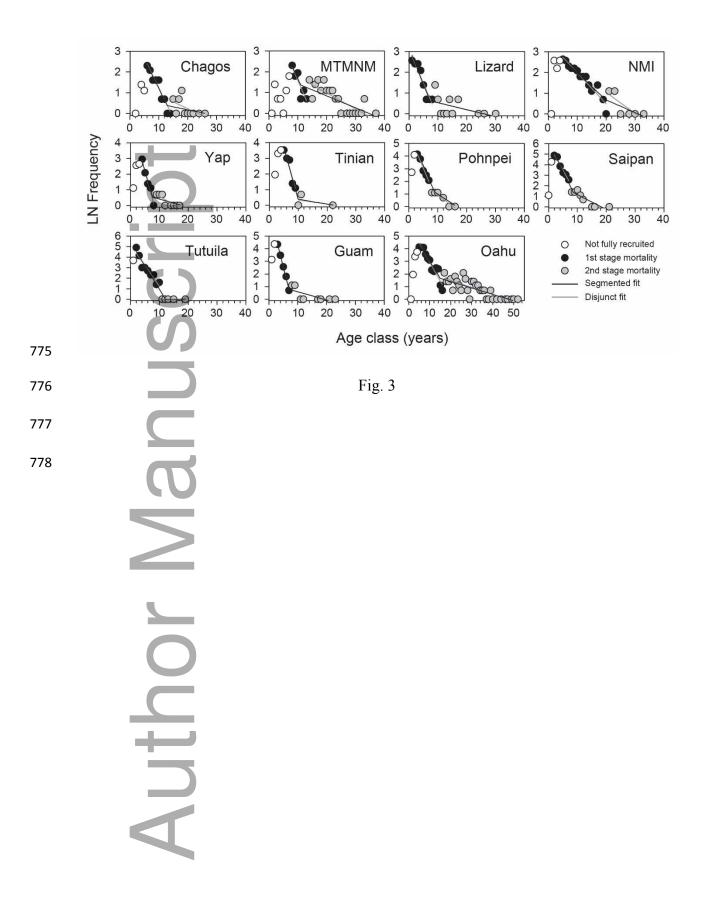
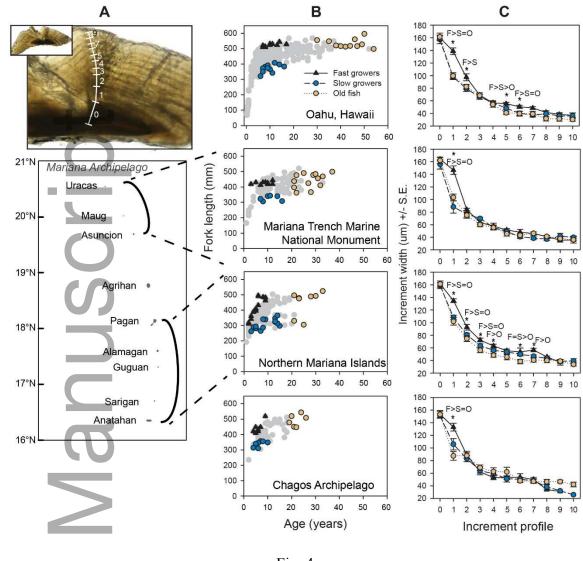


Fig. 2

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Fig. 4

