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*CORRESPONDENCE Kim Bassos-Hull kbhull@mote.org

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Whitespotted eagle ray (*Aetobatus narinari*) age and growth in wild (*in situ*) versus aquarium-housed (*ex situ*) individuals: Implications for conservation and management

Atlantine Boggio-Pasqua¹, Kim Bassos-Hull^{2,3*}, William H. Aeberhard⁴, Lisa A. Hoopes⁵, Denise A. Swider⁶, Krystan A. Wilkinson^{2,3} and Manuel Dureuil⁷

¹Aix-Marseille Uni, Marseille, France, ²Sharks and Rays Conservation Research Program, Mote Marine Laboratory, Sarasota, FL, United States, ³Chicago Zoological Society's Sarasota Dolphin Research Program, c/o Mote Marine Laboratory, Sarasota, FL, United States, ⁴Swiss Data Science Center, ETH Zürich, Zürich, Switzerland, ⁵Research and Conservation Department, Georgia Aquarium, Atlanta, GA, United States, ⁶Discovery Cove, Orlando, FL, United States, ⁷Shark and ray conservation centre (SHARCC), Gaetz Brook, NS, Canada

Elasmobranchs typically display slow growth, late maturity and low fecundity life history characteristics, making them vulnerable to fishing pressures and environmental perturbations. The whitespotted eagle ray (Aetobatus narinari), a large pelagic migratory ray with an endangered status on the IUCN Red List, fits this pattern based on available literature. Historically, age and growth parameters for this ray have been reported through vertebral ageing methods. However, the periodicity of vertebrate band pair formation, which is used for ageing, has not been validated for this species, making ageing accuracy and thus the resulting growth parameters uncertain. In this study, we used both a frequentist and Bayesian method of estimating sex-specific von Bertalanffy growth parameters (DW_{∞}, k) in wild recaptured versus aquariumhoused rays. Additionally, we estimated growth from repeated measurement data collected from aquarium-housed rays, as an alternative approach to obtain growth parameters while allowing for individual variability. Between 2009 and 2020, 589 whitespotted eagle rays were caught, measured, tagged and released along the southwest coast of Florida. Of these rays, 34 were recaptured between 5-1413 days at liberty. Nineteen additional rays were collected during the same period, transported and maintained at Georgia Aquarium, Atlanta, where they were regularly weighted and measured. Data from Association of Zoos and Aquariums accredited facilities provided prior information on maximum size for the Bayesian estimations, and size at birth, size at maturity, and maximum life span. These data were used to plot and interpret von Bertalanffy growth curves. Wild whitespotted eagle rays were found to grow faster and mature earlier than previously thought, with Bayesian estimates of k = 0.28 year⁻¹ in females, and k = 0.30 year⁻¹ in males. Aquariumhoused individuals seemed to grow slower and reach smaller sizes, although data provided by the aquariums showed variable growth patterns depending on the facility. Longevity was estimated at 14-15 years in wild rays while maximum lifespan observed in aquariums was 19-20+ years. Life history parameters and growth trajectories generated from this study offer valuable information to aid with future conservation management strategies of this endangered species.

KEYWORDS

eagle ray, life history, aquarium, tag-recapture, age and growth analysis, maturity, longevity

Introduction

Life history traits (e.g., growth patterns, age at maturity or longevity) determine how populations will change in time and respond to exploitation pressure, making them key aspects in species management and conservation. Many elasmobranch (shark, ray and skate) species are particularly vulnerable to overexploitation (Frisk et al., 2005; Dulvy et al., 2014) and are among the latest-maturing and longest-living vertebrates (Hoenig & Gruber, 1990; Cortés, 1998; Cortés, 2002). Estimating age at maturity or longevity requires accurate age information, which is often difficult to obtain reliably in sharks and rays. The most widely used ageing technique in elasmobranchs consists of counting growth zones in calcified structures, such as band pairs in vertebral centra, assuming periodic deposition (Cailliet & Goldman, 2004). However, only a few studies have rigorously validated this temporal periodicity for particular species and commonly only for some portion of lifespan (see review by Cailliet, 2015). To further complicate this, recent studies suggest that growth zone deposition (i.e., vertebral band pairs) in sharks and batoids is closely related to somatic growth and body shape, rather than time (Natanson et al., 2018; James & Natanson, 2020). As an alternative, theoretical ages may be obtained indirectly from length via growth models, such as the von Bertalanffy growth model, which is the most common growth model utilized for fishes. Mark-recapture data can provide information on size increments over time, which are used as inputs to these growth models. Rearing in research laboratories or public aquariums such as Association of Zoos and Aquariums (AZA) accredited institutions also provides the invaluable opportunity to advance scientific knowledge of the animals in their care and enhance the conservation of wild populations (Association of Zoos and Aquariums, 2022a). Carefully and regularly taken morphometric data from participating aquariums can help fill critical data gaps in our understanding of age and growth, especially of shark and ray species for which band pair formation periodicity has been difficult to validate.

The whitespotted eagle ray is a durophagous mesopredator feeding mainly on benthic invertebrates (Ajemian et al., 2012; Serrano-Flores et al., 2018) and is often found near coral reefs, along beaches and coastal inlets, and in estuarine seagrass habitats (Silliman & Gruber, 1999; Ajemian et al., 2012; Bassos-Hull et al., 2014; Flowers et al., 2017; Cerutti-Perevra et al., 2018; DeGroot et al., 2020). Previously thought to be a single, globally distributed species in warm and tropical waters (Bigelow and Schroeder, 1953; Last et al., 2016) and the only species of its genus, A. narinari has been recently identified as a complex of several cryptic lineages: the genus Aetobatus (Blainville, 1816) currently comprises at least five known species, based on genetic and morphologic evidence (Richards et al., 2009; White, 2014; White & Naylor, 2016; Sales et al., 2019), with the "true" A. narinari now restricted to the tropical Atlantic Ocean (Sales et al., 2019; Fricke et al., 2022). The conservation status of each member of this species complex thus needs to be assessed accounting for their geographic range and regionalized threats. While A. narinari is protected in Florida state waters, Bermuda and the Bahamas, this protection might be insufficient as the species is capable of long-distance migrations over hundreds of kilometers (see Sellas et al., 2015, DeGroot et al., 2021 and Friess et al., 2021) and is exposed to intensive and unregulated inshore fisheries throughout most of its range, including for consumption in parts of Central and South America (Cuevas-Zimbrón et al., 2011; Tagliafico et al., 2012; Cordovés et al., 2013; Cuevas et al., 2013; Cuban Ministry of the Food Industry, 2015). Females bear one to five pups after a gestation period of approximately one year (Swider et al., 2017). Due to its low fecundity, association with near-shore coral reef, beach, and seagrass habitats, and its exposure to fishing pressure, A. narinari was recently classified as "Endangered" by the IUCN Red List (Dulvy et al., 2021).

Current knowledge on *A. narinari* life history (Table 1) is limited and derived from a few fisheries-dependent studies (Dubick, 2000; Yokota & Lessa, 2006; Tagliafico et al., 2012; Utrera-López, 2015; Araújo, 2020) and one tagging study (Bassos-Hull et al., 2014). Dubick (2000), Utrera-López (2015)

and Araújo (2020) used calcified vertebrae to visualize band pairs and estimate growth parameters in Puerto Rico, Mexico and Brazil respectively. However, they were not able to reliably validate the band pair formation periodicity. Bassos-Hull et al. (2014) was the first tagging study on wild, free-swimming A. narinari in US coastal waters, investigating life history and seasonality of the species along the southwest Florida coast (i.e., Gulf of Mexico). Using the Fabens method (Fabens, 1965), von Bertalanffy growth parameters were estimated from recaptured rays' disc width increments during time at liberty, but the sample size (n=22) was small and included some biologically questionable (e.g., null and negative) growth rates. As for ex situ studies, little research has been done on A. narinari in managed care due to the relatively few individuals represented in zoo and aquarium collections. Although there are only 53 A. narinari currently housed in 16 different AZA-accredited facilities (Swider et al., 2021), the species is becoming increasingly common in public aquariums due to their visually appealing spot pattern and graceful swimming motions (Swider et al., 2017).

The typically small sample sizes available for both wild and aquarium populations, combined with the possibility of measurement errors and non-negligible individual growth variability, imply that simple growth estimation methods such as the (frequentist) Fabens method can yield unrealistic growth parameters (see Bassos-Hull et al., 2014). Incorporating auxiliary information on growth parameters, even basic information such as a biologically feasible range for the disc width, can help the estimation process in such situations (e.g. Dureuil et al., 2022). Building on previously published work by Bassos-Hull et al. (2014) and incorporating data from individuals housed in aquarium and zoo facilities, this study re-estimated growth parameters in this endangered ray species using both a frequentist and Bayesian modeling framework. In the latter, auxiliary information was formalized in prior distributions assigned to growth parameters. In addition, important life history parameters such as size at birth, age at maturity, growth, maximum size and longevity of wild and aquarium-housed A. narinari were examined and compared. We propose that using a combination of information obtained through in situ and ex situ studies will improve these life history and growth parameter estimates, which is critically important to stock assessments and management agencies and will help aquarium facilities enhance their husbandry methods for the species.

Materials and methods

The study area, the sampling process, and the animal captures and handling for the wild rays are exactly the same as described in Bassos-Hull et al. (2014), except only boat-based surveys are considered in this study. The following two

paragraphs summarize the adopted field methodology. Please refer to the Methods section of Bassos-Hull et al. (2014) for further detailed information.

Study area and sampling surveys

A. narinari individuals were captured in the eastern Gulf of Mexico ("Gulf") through boat-based surveys (n=321) conducted between July 2009 and November 2019 along the southwest Florida coast (Figure 1A). This area consists mostly of fringing barrier islands and shallow passes and inlets (200–2,500 m wide, < 20 m deep), which shelter various invertebrates that are potential prey for *A. narinari*: macrogastropods such as whelks and conch, and bivalves such as scallops and clams (Ajemian et al., 2012; Serrano-Flores et al., 2018). Sampling occurred primarily between north Longboat Key (latitude 27.4°N, longitude –82.7°W) and south Siesta Key (latitude 27.2°N, longitude –82.5°W) (Figure 1B), predominantly April through November each year.

Animal captures, tagging and measurement

When rays were encountered in workable conditions of depth and current, they were captured with a nylon seine net for medium to large rays or a cast net for small rays (<80 cm disc width, DW), with in-water assistance provided by one or two snorkelers to transfer them to the vessel's live well. Very large rays (>180 cm DW) were examined in a floating net pen (2.5 m diameter) off the side of the boat. Once a ray was restrained, a series of measurements and samples were taken. Males were identified by the presence of claspers, which were classified as noncalcified (soft and flexible claspers, immature), partially calcified (harder but partially flexible claspers, maturing), or fully calcified (large and rigid claspers, mature). DW and total length (TL) were measured to the nearest mm, and WT was measured to the nearest 0.1 kg using a hoop net attached to a calibrated digital scale on a davit. Each ray was injected with a passive integrated transponder (PIT), unless a previously applied PIT tag was detected; in this case it was used to identify the individual and the event was recorded as a recapture. All captures and recaptures were performed by trained researchers following this protocol.

A subset (n=19) of the animals captured for the wild study were retained for eventual display at Georgia Aquarium (GAI). Three of these individuals were collected in March 2009 prior to the initiation of the field research study and 16 were collected between 2012-2015 during the field research study. All animals collected and sampled for projects were conducted under Florida Protected Species permit #s SAL-(09-19)-1140-SRP approved protocols. All animal handling procedures were approved



through Mote Marine Laboratory's IACUC permits #10-03-PH1 and 13-02-PH1.

measuring tape. WT was recorded to the nearest 0.1 kg using a hanging scale. After examination, *A. narinari* individuals were released into the main exhibit and resumed normal behavior.

Animal housing and monitoring in the study aquarium

Current and historical morphometric data from these 19 aquarium-housed A. narinari were included in this study. All A. narinari at GAI were housed in the Ocean Voyager exhibit, in a 24-million-liter saltwater habitat containing about 8000 individuals of 60 other species of sharks, rays, and teleost fishes. Water temperature in the exhibit was maintained between 24-25°C year-round. Juvenile and adult rays were target fed a daily ration between 1.5-3% of their body weight (BW), with pups receiving higher rations (3-5% of BW/d). Current diets consist of surf clam (Spisula solidissima), hardshell clam (Mercenaria mercenaria), Jonah crab (Cancer borealis), knobbed whelk (Busycon carica), Atlantic sea scallops (Placopecten magellanicus), whiteleg shrimp (Litopenaeus vannamei) and blue mussels (Mytilus edulis), but have varied historically. Animals were handled at least once a year for routine veterinary examination including morphometrics (DW and WT) and more frequently as veterinary or husbandry needs dictated. For these examinations, animals were captured by divers and transferred from the main exhibit into a small holding pool with oxygenated water. DW was measured to the nearest 0.1 cm using a soft

Aquarium survey

Aetobatus narinari is managed in public aquariums through the Species Survival Plan[®] (SSP) of the Association of Zoos and Aquariums. The cooperatively-managed SSP programs oversee the population management and enhance conservation of select species in AZA member institutions. SSP programs are led by AZA member volunteers who work collaboratively with AZA committees, Scientific Advisory Groups, and AZA-accredited institutions to maintain healthy, genetically diverse, and demographically varied populations through strategic management and planning (Association of Zoos and Aquariums, 2022b). To collect data for this study, a survey was sent to the AZA A. narinari stakeholders. In the survey, institutions were asked to provide the following information, as available, regarding animals of each sex in their collection: maximum size (DW in cm) recorded with corresponding age, maximum lifespan recorded and corresponding size, all sizes at birth recorded and ages at first reproduction recorded for sexually mature rays. Participants were asked to indicate how age was determined, e.g. based on the size at capture and the time spent in the aquarium facility, or exact age if the ray was born in an ex situ setting.

Growth analysis

As basis for our growth model we considered the von Bertalanffy (vB) equation (von Bertalanffy, 1938):

$$DW_t = DW_{\infty} - (DW_{\infty} - DW_0)e^{-kt} \tag{1}$$

where DW_t is the length at age t, DW_0 is the length at birth (t = 0), DW_{∞} is a parameter representing the (asymptotic) maximum length expressed in the same units as DW_t and, DW_0 and k is a parameter known as the Brody coefficient describing how fast DWgets to DW_{∞} as t nears ∞ , expressed in the reciprocal of the time units (e.g., year⁻¹). The original vB equation implicitly takes birth as a reference point and defines a growth increment from t = 0 to an arbitrary age t (one can subtract DW_0 on both sides of Equation (1) to make this apparent). A more general form of the vB equation is given by Fabens (1965) for two arbitrary time points T_1 and T_2 , with $T_1 \leq T_2$, and the corresponding lengths DW_{T_1} and DW_{T_2}

$$DW_{T_2} = DW_{\infty} - (DW_{\infty} - DW_{T_1})e^{-k\Delta T}$$
(2)

where $\Delta T = T_2 - T_1$. This formulation is therefore suitable to examine growth from the present *A. narinari* data. T_1 then denotes the time of first capture and T_2 is the time at recapture. The absolute times themselves do not matter, only the time difference ΔT (i.e., the time at liberty) enters the growth model.

Equation (2) represents merely an idealized growth trajectory. As such, it can only be expected to hold on average for growth measurements, assumed collected with some random errors. This yields the following Fabens (1965) equation formulated as a statistical model (see e.g. James, 1991), which we will simply refer to as the Fabens model:

$$DW_{T_2,i} = E[DW_{T_2,i}] + \epsilon_i$$

= $DW_{\infty} - (DW_{\infty} - DW_{T_i,i})e^{-k\Delta T_i} + \epsilon_i,$ (3)

where the *i* index identifies an individual among *n* in a given sample (*n* being the sample size), $E[DW_{T_{2'}i}]$ denotes the mean of a random variable (mathematical expectation), and ϵ_i is an error term with mean 0 and a constant variance denoted by σ^2 . In addition, the error terms are assumed independent. This means that the way the measurement $DW_{T_{2'}i}$ differs from its mean E $[DW_{T_{2'}i}]$ cannot be predicted from how the length measurement of another individual measurement $DW_{T_{2'}j}$ differs from its respective $E[DW_{T_{2'}j}]$, for $i \neq j$. Under these assumptions, Equation (3) closely resembles to a non-linear regression model where the lengths at recapture $DW_{T_{2'}i}$ form the response variable while the lengths at capture $DW_{T_{2'}i}$ and the times at liberty ΔT_i are covariates (explanatory variables) considered fixed. The unknown parameters to estimate are DW_{∞} , k, and the error variance σ^2 .

Frequentist estimation of DW_{∞} and k can be carried out by least squares, i.e., the estimates $\widehat{DW_{\infty}}$ and \hat{k} jointly minimize the sum of squared residuals which is equivalent to maximizing a

likelihood function based on assuming all ϵ_i 's are identically distributed as Gaussian. No closed form expressions exist (Fabens, 1965), these have to be found numerically for instance through a Newton-Raphson algorithm. Given these estimates, the error variance σ^2 is typically estimated from the mean squared residuals corrected for the lost degrees of freedom:

$$\hat{\sigma}^{2} = \frac{1}{n-2} \sum_{i=1}^{n} (DW_{T_{2},i}(\widehat{DW_{\infty}} - (\widehat{DW_{\infty}} - DW_{T_{1},i})e^{-\hat{k}\Delta T_{i}}))^{2}$$

The maximum likelihood estimator (MLE) for σ^2 is given by $\hat{\sigma}^2(n-2)/n$. Standard errors can then be computed from the asymptotic approximation given by the inverse Fisher information matrix. The latter also indicates that \widehat{DW}_{∞} and \hat{k} are always negatively correlated.

A likelihood ratio test (LRT) was constructed to compare growth parameters between wild and aquarium rays. LRT pvalues were computed two different ways, using the asymptotic χ^2 distribution and through a parametric bootstrap scheme (see Supplementary Material 1 for detailed methodology).

In addition to the frequentist estimation and inference described above, we also applied a Bayesian approach. We refer to Gelman et al. (2014) for a general background on Bayesian statistics. The main addition here is that the data analyst incorporates subjective knowledge and uncertainty about the Fabens model parameters $\mathit{DW}_{\scriptscriptstyle{\infty}}$, k, σ^2 and in the form of probability distributions that do not depend on the data at hand, referred to as prior distributions. Following Dureuil et al. (2022), we constructed a lognormal prior for DW_{∞} based on published studies (Silliman and Gruber, 1999; Dubick, 2000; Cuevas-Zimbrón et al., 2011; Ajemian et al., 2012; Tagliafico et al., 2012; Utrera-López, 2015; Briones Bell-lloch, 2016; DeGroot et al., 2021; Araújo et al., 2022) and on our aquarium survey results for the wild and aquarium populations, respectively. For both populations, we used the largest reported length DWmax, for both females and males, to set the lognormal prior mean and variance (on the natural logarithm scale) parameters: the lognormal median is set as DWmax / 0.99 so that the mean parameter is $\log (DW_{max}/0.99)$ while, given the mean, the variance is found numerically such that the 99th lognormal percentile matches 1.2 times the median. The 1.2 coefficient ensures a reasonably wide distribution with some inherent right-skewness that is meaningful for an asymptotic parameter like DW_{∞} for which more uncertainty exists for higher values than for smaller ones; see Supplementary Material 2, for the lognormal prior distributions we thus specified for wild and aquarium, and for female and male, A. narinari individuals. Regarding the other two parameters, k and σ^2 , we had little prior information which could be directly represented by a probability distribution. We followed Dureuil et al. (2022) in specifying uniform prior distributions for both parameters, with lower and upper bounds defining a realistic range of values: [0.01, 1] for k and $[10^{-5}, 50]$ for σ . These bounds

are the same for females and males, as well as for aquarium and wild individuals.

Based on our priors and a Gaussian distribution assumed for the ϵ_i error terms, i.e., a Gaussian likelihood for the data given the parameters, Bayesian estimates are based on the posterior distribution of the parameters given the data. This posterior distribution is not available in a closed form here, thus we relied on the No-U-Turn Sampling (NUTS) algorithm (Hoffman and Gelman, 2014) which can be considered as an advanced Markov chain Monte Carlo (MCMC) method. Running five chains in parallel, we retained the last 10,000 draws after a burn-in period of 10,000 iterations. This means we had 50,000 draws from the posterior distributions from which we derived summary statistics and graphical representations like histograms (see Supplementary Figures 3.1-3.4). To ease the comparison of estimates between frequentist and Bayesian estimation, we considered a Bayesian point estimate computed as the median of the posterior draws. Using a posterior median rather than, say, a posterior mean, is meaningful here due to the high rightskewness we regularly notice in the posterior distributions, notably in that of DW_{∞} with small sample sizes. We computed credible intervals following the highest density interval approach; a 95% credible interval consists of the narrowest interval which contains 0.95 probability around the posterior mode.

A formal comparison of growth parameters between wild versus aquarium populations was carried out by computing Bayes factors (BFs; Kass & Raftery, 1995). The exact methodology is available in Supplementary Material 4.

The wild versus aquarium population comparison requires the same model to be fitted to both samples, which is the Fabens model in Equation (3) above with lengths at a single recapture modeled given the lengths measured at the initial capture. But each individual of our aquarium population was measured multiple times, at their capture and throughout their lives at the aquarium. This extra information was leveraged in a secondary analysis: having repeated measurements for each aquarium individual allowed us to specify a random effect, say on DW_{∞} , which represents individual growth variability. The Fabens model with random intercept on DW_{∞} is thus:

$$DW_{T_{i,i}} = DW_{\infty,i} - (DW_{\infty,i} - DW_{T_{i,i}})e^{-k\Delta T_{j,i}} + \epsilon_{i,i}$$
(4)

where $j = 2,3,...,n_i$ identifies the recapture among the n_i repeated measurements for individual $i, \Delta T_{j,i} = T_j - T_1$ for individual i, the $\epsilon_{i,j}$ error terms are independent Gaussian with mean zero and variance σ^2 , and the $DW_{\infty,i}$'s are now random too with a Gaussian distribution with mean μ_{∞} and variance σ^2_{∞} . As compared to the previous Fabens model, this model has one extra parameter: the random intercept variance σ^2_{∞} which represents how different the individual-specific vB curves are. Thus, a σ^2_{∞} value close to zero means the individuals are relatively similar to each other in terms of growth trajectories, while a large σ_{∞}^2 value (relative to σ^2) means the individual vB curves differ markedly from the population average vB curve parametrized by μ_{∞} . We fitted such a random effects model by maximizing the marginal log-likelihood of the data after having integrated out the (unobserved) $DW_{\infty,i}$'s; we again made use of the Laplace approximation for these integrals. Given parameter estimates, the random intercepts can then be predicted by the maximizers of the joint log-likelihood (equivalent to posterior modes if we were to view the random effect distribution as a Bayesian prior) as these are anyway computed in the Laplace approximation scheme.

All the model estimation and inference described above were implemented in R version 4.0.2 (R Core Team, 2022). We made use of the R package Template Model Builder (TMB; Kristensen et al., 2016) to code the joint likelihood of the Bayesian Fabens estimation and of the Fabens model with random intercept. For the former, we used the R packages tmbstan (Monnahan & Kristensen, 2018) and rstan (Stan Development Team, 2020) to link to the Stan Stan Development Team, 2022 statistical libraries which include the NUTS algorithm. We used TMB's efficient implementation of the Laplace approximation for integrating random effects both to evaluate the marginal loglikelihood in the random effects model and to evaluate the marginal densities in the BF computations. Our code is readily available on the GitHub page of the third author (Aeberhard, 2022).

Data preparation

In the wild population, only recaptures with times at liberty longer than 90 days were included in the growth analysis because of the large variability and influence on errors for shorter time periods (Simpfendorfer, 2000; McAuley et al., 2006). Successive recaptures of the same individual were merged into a single capture-recapture event to maximize the time at liberty and thus the observed growth. Two methods were then used to identify potential outliers, the influence plot (Dureuil & Worm, 2015) and the interquartile range (Dureuil et al., 2022). Any data point identified by both of these methods was considered an outlier. To estimate growth parameters in the aquarium population, we considered the first and the last measurement of each individual in order to maximize the observed growth.

Life history characteristics

Following Equation (1), growth parameters estimated by frequentist and Bayesian approaches were used to plot von Bertalanffy growth trajectories of each population. DW_0 was defined for each sex as the average size at birth based on sizes reported in the aquarium survey. Size at sexual maturity, i.e., the disc width at which 50% of the individuals are mature, was

Lifespan	Unknown, > 20-25 years (Dubick, 2000)
Maximum size (DW = disc width)	 190 cm DW for males (190 cm in Tagliafico et al., 2012; 185 cm in Bassos-Hull et al., 2014) 226 cm DW for females (231.1 cm in Bigelow & Schroeder, 1953; 226 cm in Tagliafico et al., 2012)
Size at birth	~ 40-45 cm DW (41.4-44.7 cm in Dubick, 2000; 30-40 cm (exact measurements unknown) in Yokota & Lessa, 2006; 44.5 cm in Tagliafico et al., 2012)
von Bertalanffy growth parameters	$DW_{\infty} \sim 165-245$ cm and k ~ 0.05-0.2 for females ($DW_{\infty} = 245.9$ cm, $k = 0.03$ year ⁻¹ in Dubick, 2000; $DW_{\infty} = 200.0$ cm, $k = 0.13$ year ⁻¹ in Utrera-López, 2015; $DW_{\infty} = 164.2$ cm, $k = 0.18$ year ⁻¹ in Araújo, 2020) $DW_{\infty} \sim 140-155$ cm and k ~ 0.1-0.25 for males ($DW_{\infty} = 146.5$ cm, $k = 0.09$ year ⁻¹ in Dubick, 2000; $DW_{\infty} = 140.0$ cm, $k = 0.18$ year ⁻¹ in Utrera-López, 2015; $DW_{\infty} = 153.3$ cm, $k = 0.25$ year ⁻¹ in Araújo, 2020)
Size at maturity	 115-130 cm DW for males (122 cm in Dubick, 2000; 129 cm in Tagliafico et al., 2012; 127 cm in Bassos-Hull et al., 2014; 115.6 cm in Araújo et al., 2022) 125-135 cm DW for females (124 cm in Dubick, 2000; 135 cm in Tagliafico et al., 2012; 129.4 cm in Araújo et al., 2022)
Age at maturity	Unknown, 14.2 years for males/15.4 years for females in Dubick, 2000
Litter size	1-5 pups (2-4 pups in Dubick, 2000; 1-5 pups in Tagliafico et al., 2012)

TABLE 1 Literature summary of Aetobatus narinari life history traits.

Bold values are the summary values for each life history parameter.

determined from clasper rigidity in males (see methodology of Bassos-Hull et al., 2014) and from literature (Tagliafico et al., 2012) in females. Longevity was calculated as the age at size corresponding to 99% of DW_{∞} (Dureuil et al., 2021). The disc width-weight relationship in wild versus aquarium rays was also assessed and can be reviewed in Supplementary Material 5.

Results

Morphometrics of captured and recaptured wild rays

A total of 609 individual rays (347 males, 261 females, and 1 unrecorded sex) were captured between July 2009 and January 2020. The sex ratio of our sample was significantly skewed toward males (male:female ratio = 0.75, χ^2 = 12.5, df = 1, p< 0.001). Although female maximum size (205 cm DW) and weight (119.2 kg) exceeded those of males (191.2 cm DW, 108.4 kg), there were no significant differences between average DW and WT of males vs. females (DW: t-test, t = 0.88, df = 536.5, p = 0.380; WT: t = -0.004, df = 443.56, p = 0.996). Of 346 males and 259 females measured, male DW ranged 42–191.2 cm while females ranged 41.4–205 cm. Male WT ranged 1.1–108.4 kg and females ranged 1.3–119.2 kg. The normalized size distribution of all captured rays was plotted in Figure 2.

Of the 589 A. *narinari* individuals tagged and released during the field research study (excluding the 16 rays transferred to GAI and utilized to examine growth in the aquarium), 34 (5.8%) were recaptured on one-two occasions, including 20 males and 14 females. For recaptured rays, size at capture ranged from 51 to 159 cm DW in males and 56 to 188.4 cm DW in females. Size at recapture ranged from 59.2 to 169.5 cm DW in males and 57 to 185.2 cm DW in females. Time at liberty ranged from 5 to 1413 days (mean = 316 days) in males, and 7 to 995 days (mean = 333 days) in females.

Morphometrics and growth of aquariumhoused rays

Nineteen rays, including 13 males and six females, were monitored at GAI after being captured in the study area (Supplementary Material 6). Size at first measurement ranged



Normalized size histogram of wild and aquarium-housed Aetobatus narinari. Wild sample consists of 628 measurements from 589 individuals, with 1-3 measurements per individual. Aquarium sample consists of 146 measurements from 19 individuals, with 5-11 measurements per individual. 48.3-120 cm DW in males and 64-108 cm DW in females, and size at last measurement ranged 118-143 cm DW in males and 144-162 cm DW in females. Males were monitored during 3.3 to 10.9 years and grew 11 to 81 cm, while females were monitored during 4.4 to 7.4 years and grew 40 to 97 cm.

AZA - SSP survey results

Information was provided from nine institutions representing 75 individual animals (Table 2). Due to lack of regular morphometric measurements taken on many animals managed in the SSP, not all participating facilities were able to fully provide answers to all questions with accuracy. One newborn female of 35.5 cm DW considered abnormally small its whole life was not included in the average size at birth calculation.

Estimation and comparison of growth parameters

Estimation of von Bertalanffy growth parameters in wild versus aquarium rays

In the wild group, none of the eligible data points were considered an outlier using the methods previously described (influence plot and IQR). However, a single recapture of a small male (74 cm DW) showing null growth over five months (166 days) was removed from the dataset. The selection of recaptures with times at liberty longer than 90 days resulted in a final data set of 22 recaptures (15 males and seven females) used for growth analysis (Supplementary Material 7). Size at capture

TABLE 2 Summary of the participating AZA aquariums' data.

ranged from 60 to 155 cm in females, and from 51 to 159 cm DW in males. Size at recapture ranged from 134 to 185.2 cm DW in females, and from 84 to 169.5 cm DW in males. Time at liberty ranged 1.1–2.72 years (403–995 days) in females and 0.37–3.87 years (134–1413 days) in males.

The frequentist and the Bayesian approaches produced consistent, plausible growth parameter estimates for all groups of rays, except an unlikely high frequentist estimate of *k* in the aquarium females (Table 3). We note that this large value for *k* implies a likely underestimated DW_{∞} , since these two parameter estimates are negatively correlated. This explains the discrepancy between frequentist and Bayesian estimates of DW_{∞} for the aquarium females. Females were estimated to have a larger DW_{∞} than males, this sexual dimorphism being more pronounced in the wild population ($\Delta DW_{\infty} \sim 39$ cm) than in the aquarium population ($\Delta DW_{\infty} \sim 31$ cm). *k* estimated values ranged between 0.28 and 0.32 year⁻¹ in the wild rays, while ranging 0.36–0.46 year⁻¹ in the aquarium rays.

Comparison of growth parameters between groups: Varying by sex and condition (wild/aquarium)

All the *p*-values computed using the asymptotic χ^2 distribution were well under the 0.05 threshold, meaning the growth parameters (DW_{∞}, k) were significantly different between groups in the four comparisons (Table 4). The (more reliable) bootstrapped p-values were also under the 0.05 threshold but weaker, with some values close to 0.02–0.03.

The Bayes factors only indicated substantial evidence for different growth parameters between wild and aquarium males (BF>>3.2). Simulations (not presented here), however, revealed that on average only 57% of BF values were >3.2 when the growth

	DW at birth(cm)	DW at sexual maturity(cm)	Maximum DW (cm)	Maximum observed longevity (years)
Males	39-55 (48.1 ± 4.4)	119–144	150*	19*
	(n=13)	(first successful mating)		(time spent in the facility)
Females	41-58.4 (51.5 ± 6.1)	135.3–160	186.7*	20*
	(n=8)	(first parturition)		(time spent in the facility)

*Four different individuals, still alive as of 2021.

TABLE 3 von Bertalanffy growth parameter estimates.

Group (<i>n</i>)	Frequentist Estimation			Bayesian Estimation		
	DW_{∞} (cm)	k (year ⁻¹)	σ (cm)	DW_{∞} (cm)	k (year ⁻¹)	σ (cm)
Wild females (7)	222.6	0.29	8.1	225.7 [196.0-258.4]	0.28 [0.18-0.42]	8.0 [4.2-14.7]
Wild males (15)	183.7	0.32	7.1	187.6 [169.7-209.3]	0.30 [0.20-0.42]	7.1 [4.8-10.4]
Aquarium females (6)	153.8	5.00	8.1	166.9 [150.7-197.8]	0.36 [0.12-0.92]	10.9 [4.3-23.6]
Aquarium males (13)	134.1	0.46	6.9	136.2 [128.4-148.5]	0.41 [0.18-0.87]	7.1 [4.5-10.8]

For Bayesian estimations, 95% credible intervals (highest density) from posterior draws are in brackets.

	Likel	Bayes Factor	
Group 1 (<i>n</i>) / Group 2 (<i>n</i>)	χ^2 p-value	Bootstrapped p-value	
Wild females (7) / Wild males (15)	0.0062	0.024	0.19
Aquarium females (6) / Aquarium males (13)	0.0002	0.0015	1.97
Wild females (7) / Aquarium females (6)	0.0059	0.0172	0.29
Wild males (15) / Aquarium males (13)	4.67E-07	< 1E-04	3258.67

TABLE 4 Comparison of growth parameters between groups.

parameters were indeed different (with simulation parameters set to the Bayesian estimates in Table 3 so as to mimic the real data), suggesting a lack of test power.

Estimation of individual variability in DW_{∞}

Repeated measurements of aquarium rays were used to investigate the individual variability in DW_{∞} while keeping k constant across individuals. Each ray counted 4–10 measurements along their lifespan at the GAI, 4–6 in females and 4–10 in males. We found a high variability in DW_{∞} ($\sigma_{\infty} = 30.3$ cm) for the females (Table 5). In comparison, males yielded more consistent results with a moderate variability in DW_{∞} ($\sigma_{\infty} = 11.5$ cm). Individual-specific von Bertalanffy growth curves were then plotted using these parameter estimates (Figure 3).

Growth trajectories and life history characteristics

Growth parameter estimates from Table 3 were used to plot average von Bertalanffy growth trajectories for each population (Figure 4). For plotting purposes, size at birth DW_0 is required, which was determined from the average reported in the aquarium survey: 51.5 cm in females and 48.1 cm in males (Table 2).

Size at sexual maturity (i.e., the disc width at which 50% of the males are mature) was calculated from the rigidity of claspers in the wild capture data set (Clark and von Schmidt, 1965). With 288 males examined, the logistic regression produced an estimate of 128.6 cm. The disc width at which 50% of the females are mature was estimated by Tagliafico et al. (2012) as 134.9 cm. From this, age at maturity was estimated between two and three years old in wild rays, and up to six years old in aquarium males (Table 6). Longevity estimates ranged from 10

TABLE 5 $\,$ Estimated growth parameters with individual variability in $DW_{\rm co}$

Group (n)	μ_{∞} (cm)	σ_{∞} (cm)	k (year ⁻¹)	σ (cm)
Aquarium females (6)	225.9	30.3	0.11	4.9
Aquarium males (13)	145.9	11.5	0.24	5.5

to 15 years, with credible intervals overlapping between aquarium and wild rays.

Discussion

Comparison of our results with vertebral age/growth studies

Our results suggest wild A. narinari grows considerably faster (Figure 5) than previously published (Table 1). For example, on average, a ray of 100 cm DW was previously estimated to grow 20 cm in 1.7 year (female growth in Utrera-López, 2015) to 6.4 years (male growth in Dubick, 2000), whereas such growth seems to occur in 7 to 11 months on average based on our analysis. Although inherent differences between growth curves calculated from agelength and length-increment data prevent a direct comparison from being made (Francis, 1988), growth rates estimated here substantially exceed those from previous vertebral ageing studies on the species. Only few published studies investigating age and growth in elasmobranch species have successfully validated an annual deposition periodicity in vertebral centra (reviewed by Cailliet, 2015), and while a lack of validation does not invalidate deposition periodicity in one species, unvalidated age estimates resulting from vertebral readings must be carefully considered. Recent studies have highlighted several limitations when using calcified growth zones to determine age, including systemic age underestimation in larger and older individuals (Harry, 2017), and correlation of vertebral band pairs with somatic growth rather than time (Natanson et al., 2018; James & Natanson, 2020). Further investigation is therefore needed to elucidate vertebral band pair deposition and age determination in elasmobranch species such as A. narinari, and indirect, complementary approaches such as markrecapture and aquarium rearing combined with chemical markers (e.g., oxytetracycline) will certainly help answer this critical biology question.

Life history profile and implications for species management

Size increments from mark-recaptured individuals produced plausible growth parameters with the Bayesian methodology



adopted. DW... estimates for both sexes (225.7 cm in females and 187.6 cm in males) are compatible with observed maximum sizes in the region: 195.2 cm and 140 cm in the Bahamas (Silliman & Gruber, 1999) and 202 cm and 150 cm in the southern Gulf of Mexico (Cuevas-Zimbrón et al., 2011) for females and males, respectively. The largest male observed in this study (DW = 191.2 cm) is the largest male recorded in the Atlantic Ocean. In both wild and aquarium rays, females were estimated to reach larger sizes and grow faster than males. This sexual dimorphism in A. narinari was reported in fishery-dependent studies (Cuevas-Zimbrón et al., 2011; Tagliafico et al., 2012) and is typical in myliobatiform rays (Capapé et al., 2007; Schluessel et al., 2010; Setyawan et al., 2022). Larger size of females may be attributed to the accommodation of developing young, as this feature is also common for many viviparous shark species (Cortés, 2000). k estimates (0.30 year-1 in females and 0.28

year⁻¹ in males) fall within the 0.2–0.3 range theorized for rays by Holden (1974) but are relatively high compared to other myliobatiform ray species (Table 7). However, all these studies used vertebral ageing methodology with the possible limitations mentioned previously in band pair periodicity validation; thus, the resulting growth parameters of these previous studies should be carefully considered. All other things being equal, a higher kmeans that on average, the individuals approach their DW_{∞} relatively faster, resulting in earlier age at maturity and shorter theoretical longevity. Our results suggest A. narinari matures at 2-3 years old, which is half the age previously proposed (4-6 years old) in the previous IUCN assessment (Kyne et al., 2006). However, this previous estimate was based on A. ocellatus in Australian waters (Last & Stevens, 2009) which has since been re-described as a distinct species from A. narinari (White et al., 2010) and is known to reach larger sizes (Kyne et al., 2016).



Likewise, theoretical longevity appears to be shorter than previously thought, 14–15 years instead of 20–25 years estimated by Dubick (2000). Such life history parameters might be beneficial to the species' resilience since fast-growing species are theoretically less susceptible to overfishing. Nevertheless, *A. narinari* is still characterized by a low reproductive potential with a long gestation and small litters (Swider et al., 2017).

As highlighted in the recent IUCN assessment (Dulvy et al., 2021), *A. narinari* population trends are inferred from limited landings data and remain uncertain in many parts of its range. Stock assessments ideally rely on both fisheries and biological

	Frequentist Es	timation	Bayesian Estimation		
Group (n)	Age at maturity (years)	Longevity (years)	Age at maturity (years)	Longevity (years)	
Wild females (7)	2.3	15.0	2.3 [1.8-2.8]	15.3 [9.6-22.2]	
Wild males (15)	2.8	13.4	2.9 [2.3-3.5]	14.3 [9.3-20.4]	
Aquarium females (6)	0.4*	1.0*	3.5 [1.5-6.2]	11.6 [4.2-28.6]	
Aquarium males (13)	6.0	9.0	6.0 [3.3-8.2]	10.3 [4.1-18.3]	

TABLE 6 Life history characteristics inferred from von Bertalanffy growth curves.

For Bayesian estimations, 95% (highest density) credible intervals from posterior draws are in brackets.

*These very small values come from the unlikely high k estimate (see Table 3).

data related to survival and reproduction of the species. The parameters estimated in this study (i.e., DW_{∞} , k, size at birth, age at sexual maturity and longevity) contribute to refine our current knowledge of *A. narinari* life history (Table 1) and constitute helpful data to fisheries stock assessments as they can be utilized to compute essential demographic parameters, such as natural mortality (Dureuil & Froese, 2021; Dureuil et al., 2021). As regional variability may exist between distinct populations (King & McFarlane, 2010; Bradley et al., 2017), life history parameters estimated in this study are thought to be relevant at least to the Gulf of Mexico and the Caribbean Sea regions where targeted, intensive, and unregulated inshore fisheries occur (Cuevas-Zimbrón et al., 2011; Tagliafico et al., 2012; Cuevas et al., 2013).

Differences observed between aquarium and wild rays

With Bayesian estimates of $DW_{\infty} = 166.9$ cm and $DW_{\infty} = 136.2$ cm for females and males respectively, aquarium rays in this study were characterized by smaller asymptotic sizes than wild rays ($DW_{\infty} = 225.7$ cm and $DW_{\infty} = 187.6$ cm). The complementary growth analysis using multiple measurements for each ray yielded mean DW_{∞} estimates ($\mu_{\infty} = 225.9$ cm and $\mu_{\infty} = 145.9$ cm) falling closer to those estimated in wild rays, but indicated a substantial individual variability in growth trajectory, especially in females. However, a potential limitation in this analysis lies in the *k* parameter which is assumed to be constant



FIGURE 5

Von Bertalanffy growth curves generated for wild *A. narinari*. Parameters for female curves: $DW_{\infty} = 225.7$ cm, k = 0.28 year⁻¹ (this study); $DW_{\infty} = 164.2$ cm, k = 0.18 year⁻¹ (Araújo, 2020); $DW_{\infty} = 200.0$ cm, k = 0.13 year⁻¹ (Utrera-López, 2015); $DW_{\infty} = 245.9$ cm, k = 0.03 year⁻¹ (Dubick, 2000). Parameters for male curves: $DW_{\infty} = 187.6$ cm, k = 0.30 year⁻¹ (this study); $DW_{\infty} = 153.3$ cm, k = 0.25 year⁻¹ (Araújo, 2020); $DW_{\infty} = 140.0$ cm, k = 0.18 year⁻¹ (Utrera-López, 2015); $DW_{\infty} = 146.5$ cm, k = 0.09 year⁻¹ (Dubick, 2000).

Family	Species	Reference	Area	n	DW_{∞} (cm)	k (year ⁻¹)
Aetobatidae	Aetobatus ocellatus	Schluessel (2008)	Queensland (AUS) & Taiwan	111	(Gompertz) 365.6 (Gompertz) 274.9	0.026 0.037
	Aetobatus flagellum	Yamaguchi et al. (2005)	southwestern Japan	281	152.7 131.8	0.111 0.133
Myliobatidae	Aetomylaeus bovinus	Başusta & Aslan (2018)	eastern Mediterranean (Turkey)	47	242.6 238.4 (Gompertz, sexes combined) 162.14	0.056 0.044 0.171
	Myliobatis californicus	Martin & Cailliet (1988)	Monterey Bay, California	132	158.7 100.4	0.100 0.229
Rhinopteridae	Rhinoptera bonasus	Fisher et al. (2013)	Chesapeake Bay, Virginia	537	106.3 97.1	0.193 0.274
	Rhinoptera bonasus	Neer & Thompson (2005)	northern Gulf of Mexico	227	(Gompertz, sexes combined) 110	0.133
	Rhinoptera bonasus	Smith & Merriner (1987)	Chesapeake Bay, Virginia	115	125 119.2	0.119 0.126
Mobulidae	Mobula japanica	Cuevas-Zimbrón, 2012; Pardo et al. (2016)	southeastern Baja California Peninsula, Mexico	55	(sexes combined) 233.8 (Bayesian VB, sexes combined) 299.5	0.28 0.12

TABLE 7 Von Bertalanffy	growth paramet	ers estimated in Myliobat	iformes using vertebral ageing.
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Growth model by default is the von Bertalanffy. Successive rows of parameters correspond to females and males.

across all rays, whereas DW_{∞} and k are inherently negatively correlated. Considering the high variability in the small female sample, the potential limitation of a constant k parameter with variable individual DW_{∞} , and ease of comparison between A. narinari populations, we used the Bayesian estimates to derive the von Bertalanffy growth curves and the life history traits of the aquarium population. While these suggest a slower growth trajectory compared to wild rays, information communicated by the AZA facilities indicated growth variability across the aquariums, with some individuals demonstrating similar maximum sizes and growth rates (Swider, unpublished data) to wild rays. Data from the AZA survey and GAI rays also indicated similar sizes at maturity for both sexes compared to wild rays, suggesting A. narinari maturation is dependent on size rather than age. The same observation was made by Henningsen & Leaf (2010) who studied Hypanus americanus in captivity and found sizes at maturity similar to those reported in the wild. In the present study, this means GAI rays reach maturity later than wild rays.

It is well documented that animals managed under human care, sharks and rays included, can exhibit different life history traits compared to their wild counterparts (e.g., Cailliet et al., 1992; Kusher et al., 1992; Mollet et al., 2002; Braccini et al., 2010; Ezcurra et al., 2012). Growth rates in ectotherms are highly plastic in response to changes in temperature, resource availability, or density and life history trade-offs are expected in stable versus stochastic environments (Gotthard, 2001; Audzijonyte et al., 2016). The most likely explanation for the lower growth rates in aquarium *A. narinari* in this study is the year-round lower water temperatures maintained in the GAI Ocean Voyager exhibit (24-25°C). Along the Gulf coast of Florida, *A. narinari* has been shown to use only a portion (18-34°C) of the available thermal regime

available (8-34°C), and rays were mostly observed when sea temperature was above 25°C, suggesting a preference for warm waters (DeGroot et al., 2021).

Food availability can be a limiting factor for ectotherms needing to invest energy into growth, although it should be noted that optimal growth is not the same as maximal growth (Gotthard, 2001). While consumption rates for this species are unknown in the wild, daily rations for aquarium A. narinari (1.5-5% BW/d) exceeded recommendations (0.6-0.9% BW/d for adults) based on data for other species of elasmobranchs (Janse et al., 2004). Similarly, the diets of the aquarium rays were formulated to closely approximate the nutrient composition and energy density of A. narinari prey sampled in Sarasota Bay (L. Hoopes, unpublished data). Daily, stable, year-round targeted feeding of rays at GAI suggest that food availability is not limiting adequate growth as animals are reaching maturation at similar sizes to wild cohorts. In the wild, density or animal abundance can impact resource acquisition through competition for resources, shaping growth rates among fishes (e.g., Huntsmann et al., 2021), although it is difficult to empirically identify density dependent factors as a specific drivers of growth or other life-history characteristics (Audzijonyte et al., 2016). Invertebrate prey availability in Sarasota Bay is abundant and individuals can consume up to 12 different species of gastropod and bivalve prey based on stomach content analysis (K. Bassos-Hull, unpublished data), making Sarasota Bay a productive seasonal habitat for foraging. A more comprehensive understanding of the diet, feeding ecology, and energetic requirements of A. narinari would help discern the role of food intake on growth.

Bayesian growth estimates suggest a decreased longevity for aquarium rays compared with wild animals, which is a direct

result of lower asymptotic sizes. However, longevity estimates based on the von Bertalanffy parameters reflect a mean maximum age in the population, and here credible intervals considerably overlap between aquarium and wild rays (Table 6). Some studies, both in wild and aquarium contexts, have reported older rays than the longevity estimated from their von Bertalanffy parameters (Henningsen & Leaf, 2010; Vaudo et al., 2018). Likewise, several A. narinari managed under human care have lived upwards of 20 years of age (Table 2; D. Swider, personal communication), even exceeding the longevity estimated in the wild. Many animals, including elasmobranchs, have longer lifespans under human care due to readily available veterinary care, a constant supply of quality food, and freedom from predators, diseases and other pressures (Tidière et al., 2016; Grassmann et al., 2017). Conversely, elasmobranchs may also be exposed to a variety of stress-inducing variables in aquaria which may affect their health. A. narinari populations in aquarium settings have been plagued by gill parasites which can cause premature mortality (Nolan et al., 2016). As aquarium facilities perfect ideal habitat conditions and successful reproduction continues to occur in A. narinari housed in aquaria, more definitive maximum longevity for this species may be determined in the future.

Potential limitations in this study

The methods and data we used in this study have some limitations. We filtered out observations with times at liberty smaller than 90 days and one ray which displayed no growth in five months. This was out of concern of the undue impact on the estimations such zero, or even potential negative, observed growth data points can have. These would indeed bias the fitted von Bertalanffy curves downwards, especially given the small sample sizes. This filtration was a simple device to somewhat address this robustness issue. Methods which can accommodate such observations should be explored in future work.

More importantly, the estimated growth trajectories and population comparisons rely on the validity of the von Bertalanffy model itself. While we did not find any particular departures from this assumed model in our data, other more flexible models such as Gompertz or Richards growth functions (Richards, 1959; Baker et al., 1991) would be worth exploring in future analyses. Another important limitation is the Gaussian distribution assumption for all error terms (likelihood) and for the random effect on DW_{∞} for aquarium rays. Admittedly, such an assumption is often made out of convenience for ease of implementation, computation, and interpretation. However, we found model residuals to be roughly symmetrically distributed around 0 with no concerning values, and the Gaussian random effect on DW_{∞} does yield individual-specific von Bertalanffy curves which cover well the range of observed lengths (Figure 2).

Perhaps the most important limitation in this study lies in the small sample sizes. This affects both frequentist and Bayesian approaches, albeit in different ways. For the former, as noted by Dureuil et al. (2022), maximum likelihood estimates can become unreliable with extremely small samples. This is likely the case with the unrealistically large *k* we obtained for aquarium females (with n=6). Another direct impact of small sample sizes is a potentially low test power when comparing populations. Here, all LRTs pointed to significant differences, even when using a parametric bootstrap for computing p-values. Thus, we have some confidence that differences at the population level are strong enough to be reflected in our small samples here. For the Bayesian approach, the impact of the sample size is to be balanced with the amount of information supplied by the prior distributions. The specification of meaningful priors is always delicate, but even more so when the sample is small as there is a risk that the prior may be too concentrated relative to the likelihood and thus may contribute to estimation and inference more than intended. Even though our prior distributions are somewhat informative, we believe we avoided such a problem. The AZA-SSP survey supplied reliable information, in particular about the maximum observed lengths, on which we based our priors following Dureuil et al. (2022). The resulting lognormal distributions for DW_{∞} cover a realistic range of values while being adequately wide given our uncertainty. In addition, the comparison of prior densities and posterior histograms (Supplementary Figures 3.1-3.4) confirms that our priors generally do not drive estimation and inference too much. A good illustration of this sample-prior balance is the realistic k Bayesian estimate we obtained for aquarium females, where frequentist estimation seemed to break down as noted above. This sample-prior balance does however come with drawbacks, here manifested in the BF values (Table 4). With priors specified under equality of growth parameters (model M_0) reasonably wide to reflect our uncertainty, for all comparisons except the wild versus aquarium males, there is likely not enough evidence in the data, with such small sample sizes, to warrant the more complex model M_1 . In other words, when comparing with the LRT values, the differences between populations appear not strong enough to counteract the uncertainty embedded in our priors in M_0 and M_1 . The fact that the only large BF value happens for the wild versus aquarium males comparison, where the group sizes are largest here, hints that this is most likely an issue of low test power.

Conclusion

Globally, the spotted eagle ray species complex (White, 2014) have come under increased threats from targeted

fisheries and bycatch as well as habitat loss and range reduction which resulted in upgrading their IUCN Red List status, in the case of A. narinari, to "Endangered" (Dulvy et al., 2021). The evaluation assessments to categorize the IUCN Red List threat level requires reliable life history metrics such as age at sexual maturity and longevity which we were able to present. A. narinari likely grows faster and matures earlier than previously thought when comparing mark-recapture growth to vertebral banding estimates. While this may benefit the species from a fishery take perspective, other parameters such as longevity are more complicated to measure. Longevity estimates can take decades to evaluate and highlights one benefit of long term tagging studies and monitoring animals in aquariums to arrive at these estimates. The research presented in this study highlights the positive outcomes of collaborating with aquariums and how comparing in situ and ex situ individuals benefits conservation research.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding author.

Ethics statement

The animal study was reviewed and approved by Florida state laws and regulations for work on protected marine species (FWC SAL-(09-19)-1140-SRP). IACUC protocols were approved by Mote Marine Laboratory (#10-03-PH1 and #13-02-PH1).

Author contributions

KB-H conceived the study. KB-H and LH obtained funding. KB-H and KW conducted fieldwork. LH and DS provided aquarium data. AB-P, WA, KW, and MD designed the analysis methodology. AB-P, WA, and KW conducted analysis. AB-P, KB-H, WA, LH, DS, and KW helped to write the manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/ fmars.2022.960822/full#supplementary-material

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