Comparative physiological energetics of Mediterranean and North Atlantic loggerhead turtles

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Abstract

Population of loggerhead turtles nesting in the Mediterranean Sea has probably evolved from the North Atlantic (NA) population, but is geographically and genetically distinct. We aggregated previously published and new unpublished data, and took two approaches to comparing these populations: an empirical one based on statistical analyses of morphological data, and a physiological one based on a Dynamic Energy Budget (DEB) model. We then analyzed causes of faster growth and maturation, but smaller size at puberty and ultimate size of the Mediterranean (MED) loggerhead turtles relative to their NA conspecifics. The empirical analysis showed that MED eggs, hatchlings, and nesting adults were consistently smaller in terms of length and mass. The physiological approach suggested physiological adaptations of the MED population to higher salinity and scarcer food availability. In particular, these adaptations included an increase in somatic maintenance needs, and a decrease in energy investment to reach and maintain sexual maturity. Our study therefore offers a mechanistic underpinning of previously observed but unexplained life-history traits, and showcases an application of DEB theory as a tool for comparative analysis of two distinct populations of the same species.

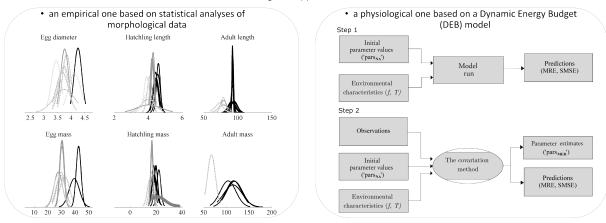
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Graphical abstract:



We compared life history traits and energy budgets of Mediterranean and North Atlantic loggerhead turtles, using two approaches:

1 1. Introduction

Loggerhead turtles inhabiting the Mediterranean Sea have probably evolved from the 2 North Atlantic loggerhead turtles more than 10000 years ago [1]. They are considered the 3 same species even though they genetically differ [2, 3], and interbreed only to a limited extent 4 when sharing the same feeding grounds [4]. The individuals from the two populations are 5 morphologically similar, but Mediterranean (MED) loggerhead turtles are smaller than their 6 North Atlantic (NA) relatives [1, 5, 6]. Nesting MED females are possibly also younger 7 than the NA ones [3, 6, 7], with shorter remigration intervals and fewer clutches per season 8 [8, 9, 10], but often more eggs per clutch [1, 5, 11]. 9 The observed variabilities in size and reproductive output could be caused by a number 10

¹¹ of generally recognized factors, notably:

(i) higher food abundance – a characteristic of the North Atlantic – results in faster growth
 rates and larger sizes (e.g., 12, 13, 14), while energy limited environments such as the
 Mediterranean reward earlier maturation at smaller sizes [5, 15];

(ii) stable conditions – a characteristic of the Mediterranean Sea – support a longer repro ductive season and higher population densities, resulting in smaller individuals due to
 less resources per individual (16, p. 297);

- ¹⁸ (iii) genetic differences cause different growth and maturation potentials [17];
- (iv) some ecological pressures (e.g., trans-oceanic migrations) favor larger sizes, while other
- ²⁰ pressures (e.g., higher adult predation) favor earlier reproduction at smaller size [5];

(v) adaptations in feeding behavior may result in different ecological niches [13].

Investigating complex interactions resulting from the aforementioned environmental and physiological factors is complicated by the large number of possible combinations, and is further hindered by a long life-cycle of loggerhead turtles encompassing several decades and multiple habitat types. Therefore, a systemic approach combining empirical data and computational analysis may help.

We undertook such an approach using previously published and unpublished data. We 27 organized our analyses in two stages. First, we performed statistical analyses based on em-28 pirical data on morphology of individuals from the two populations. Second, we developed 29 a mechanistic physiological Dynamic Energy Budget (DEB) [16, 18] model specific to the 30 Mediterranean population, and compared it to an analogous DEB model for NA logger-31 head turtles that had been described previously [19]. This is currently the only example 32 of population-specific parameter sets in the Add-my-pet project [20, 21] and likely the first 33 application of DEB models to compare two distinct populations of the same species using 34 population-specific parameter values. The environmental characteristics of the two sea basins 35 were taken into account for the comparison. By explicitly modeling the environmental fac-36 tors, we could better understand the environmental effects and possible causes of the observed 37 differences between the populations. 38

In the following sections, we briefly present the main characteristics of the environment, and the main life history traits of loggerhead turtles. The emphasis is on the Mediterranean Sea and MED loggerhead turtles (for the NA loggerhead population, please see Marn et al. [19]). We describe the collected data and our two-step approach in the methodology section, and organize the results to maintain the distinction between the two modes of analysis. Discussion sets the results in a wider context of the possible environmental and evolutionary pressures and adaptations.

46 1.1. Environmental conditions

Compared to the North Atlantic, the Mediterranean Sea is a relatively small basin – 47 $2.5 \text{ million } \text{km}^2$ [1, 22] vs. $106.5 \text{ million } \text{km}^2$, (NOAA-facts) – and its only communication 48 with other sea basins is with the Atlantic Ocean via the narrow Strait of Gibraltar. The 49 main characteristics of the Mediterranean Sea are (adapted from [1, 22, 23, 24]): (i) High 50 level of evaporation resulting in salinity difference and water deficit; (ii) Existence of two 51 main basins, the western and the eastern, connected by a shallow Sicily Channel and the 52 narrow Messina Strait, with the eastern basin more saline and warmer than the western 53 basin; (iii) Generally oligotrophic conditions resulting in the Mediterranean being one of the 54 less productive seas; and (iv) Very large biodiversity. 55

All sea turtles in the Mediterranean are listed on the IUCN Red List of Threatened 56 Species and are protected by (inter)national conventions and laws [25, 26]. Caretta caretta 57 (loggerhead turtle) is currently the most abundant sea turtle species in the Mediterranean 58 and is one of the two sea turtle species nesting there [1, 11, 27]. MED loggerhead turtles 59 mate and nest primarily in the east Mediterranean, with major nesting sites and rookeries 60 in Greece, Cyprus, and Turkey [1]. Immature NA loggerhead turtles are encountered in the 61 western Mediterranean basin [3, 4, 28]. Compared to the salinity of the North Atlantic (35-62 36 ppt, [29, 30]), salinity of the eastern Mediterranean Sea is higher (38-39 ppt, [24]), leading 63 to a hypothesis about a "salinity barrier" experienced by NA turtles [2]. 64

⁶⁵ 1.2. Loggerhead turtle life cycle

Generally the life cycle of MED loggerhead turtles is very similar to that of their NA 66 conspecifics, but some notable differences exist. Namely, MED turtles reproduce at smaller 67 sizes, especially those nesting in Libya, Tunisia, and Turkey, and they grow to a smaller 68 size (1, 31, 32, Appendix I in Tiwari and Bjorndal [5]). Age at sexual maturity is variable, 69 with estimates between 14 to 28 years for MED turtles [3, 6, 7], and similar or higher (>35) 70 years old) estimates for NA turtles [6, 33, 34]. Individuals of both MED and NA populations 71 nest every 2–3 years, laying between 1 and 5 clutches per nesting season, but Mediterranean 72 females lay on average fewer clutches into which they often deposit more eggs [1, 5, 9, 11, 35]. 73 Incubation lasts 50–60 days with the duration of incubation being inversely proportional 74 to the incubation temperature [36, 37, 38, 39]. Hatchling size varies between different nests 75 within a population, which has generally been attributed to incubating conditions such as 76 humidity, salinity, and temperature of the sand (Stokes et al. [37], Gutzke et al. [40], Packard 77 and Packard [41], Reece et al. [42], Glen et al. [43], Bower et al. [44], but see Reid et al. 78 [38], Reece et al. [42], Ji and Brana [45], Patino-Martinez et al. [46]). On average, MED 79 hatchlings are smaller than the NA hatchlings [1, 37] 80

MED loggerhead turtles seem to grow and mature faster than NA loggerhead turtles feeding in the Mediterranean Sea [3], suggesting a certain physiological adaptation of the former. Size dimorphism between populations from the two basins was noticed in other species of sea turtles, as well as fish [13, 43, 47]. However, it is unclear whether the difference in size and, in the case of loggerhead turtles, reproductive output is also a difference in the general condition of the animal, and what are the likely physiological adaptations of loggerhead turtles to the environmental conditions of the Mediterranean Sea (see also discussion online [48]).

88 2. Methods

⁸⁹ 2.1. The empirical approach – Statistical analyses of size and condition

We comprehensively searched the literature for reports on the size of loggerhead eggs, 90 hatching turtles, and nesting turtles, both for Mediterranean (MED) and North Atlantic (NA) 91 populations. Size of an individual can be specified reasonably well by measurements of length 92 and mass. However, most informative comparisons based on size include the information on 93 age. For animals encountered in the wild, age is often unknown. To bypass the unknown 94 age, we focused on discernible life history events in order to include as much data as possible 95 into the analysis. Two such events are hatching and nesting. We prioritized first nesting 96 to sexual maturation, because the exact moment of sexual maturity is hard to define and 97 observe [19, 49]. We also took into account the average egg diameter and mass, because 98 variations in egg size may explain a large part of the size variation in hatchlings [5, 43, 45, 50]. 99 We first analyzed individual samples describing eggs, hatchlings, and nesting females. All 100 data and sources are summarized in Section 2.2.1 and Appendix A. We prioritized straight 101 carapace length (SCL) to other turtle length measurements, and we used wet weight to 102 characterize body mass. When raw data were available, we tested datasets for normality 103 using Lilliefors test. When only descriptive statistics were provided, such as sample size, 104 mean and standard deviation, or mean and range of the dataset, we assumed a normal 105 distribution. In several cases, we calculated the standard deviation of the sample using 106 reported mean, range, and sample size following Hozo et al. [51]. We analyzed means and 107 standard deviations of length and mass in each life-stage using Welch's ANOVA F-test [52] 108 and visually by plotting probability density plots. 109

We subsequently used pairwise Welch's t-test of unequal variances to compare each MED sample to each of the NA samples of the same type, such that, e.g., each MED hatchling length sample was tested against each of the NA hatchling length samples. We expected NA loggerhead turtles to be larger than the MED ones, thus performing a one-sided test. To minimize the chance of false positives, we applied the Bonferroni correction (http:// mathworld.wolfram.com/BonferroniCorrection.html).

Finally, we calculated Fulton's condition index, K, as the ratio of the mass and length up cubed in g/cm³ [53] as:

$$K = W/L^3. \tag{1}$$

Fulton's condition index is often used as an indication of the physiological condition of the animal: a larger K means that the animal is better fed, i.e., has more energy available for various processes. Although originally developed for fish, K has already been applied to many species of vertebrates, including sea turtles (see Stevenson and Woods Jr [54] for an ¹²² overview), and has the advantage of not assuming a "standard" or "healthy" value that, e.g., ¹²³ relative mass indices assume [54].

To obtain the mean values of K, we pooled all samples that pertain to the same size measurement of the same life-stage and the same population. Each pooled sample was described by a mean of means (Welch's mean computed by taking into account the size and standard deviation of the sample [52]), and by pooled standard deviation [55]. We calculated the "condition index" of the egg, akin to Fulton's condition index, as the ratio of egg mass (in g) to cubed egg diameter (in cm³). We performed all analyses in Matlab R2015a.

130 2.2. The physiological approach – a mechanistic Dynamic Energy Budget (DEB) model

Life-history traits of loggerhead turtles are described well by the standard DEB model [19]. Accordingly, we characterized an individual MED turtle with three state variables: structural length (L), reserve energy (E), and maturity level (E_H) . Changes in these three state variables are due to ingestion and assimilation of energy, which is subsequently used for maintenance, growth, maturation, and reproduction. A detailed specification of the standard DEB model is given in Appendix B.

¹³⁷ We converted the state variable of the standard DEB model into measurable quantities ¹³⁸ (e.g., body length and mass) by means of conversion parameters. For example, we obtained ¹³⁹ straight (SCL) and curved (CCL) carapace lengths from structural length using two shape ¹⁴⁰ parameters, δ_{SCL} and δ_{CCL} , such that:

$$L_{\rm SCL} = L/\delta_{\rm SCL}, \ L_{\rm CCL} = L/\delta_{\rm CCL}.$$
⁽²⁾

By keeping the two shape parameters constant, we implicitly assumed isomorphic growth,thus ignoring a minor change in shape between hatchlings and late juveniles [56].

Converting the state variables to body mass, W, required including the contributions of both structure and reserve. The former contribution is by definition ρL^3 , with the density of structure being approximated by $\rho = 1 \text{ g cm}^{-3}$. The latter contribution is also proportional to ρL^3 , but only after correcting for the reserve capacity and composition via dimensionless parameter ω , and the state of reserve via food availability f:

$$W = \rho L^3 (1 + f\omega). \tag{3}$$

Quantity f, also called the scaled functional response, accounts for food availability on a scale of 0 for no food to 1 for abundant food (see Kooijman [16] for details and Marn et al. [14]).

151 2.2.1. Constructing a DEB model for Mediterranean loggerhead turtles

Parameters of every DEB model are specific to each individual. Generally, we assume 152 that similarities between individuals of the same species are greater than those between indi-153 viduals of different species. This assumption makes it possible to use data from individuals 154 of one species to estimate parameters representative of the whole species [16, 21, 57]. Here, 155 however, we extended this assumption to population-level similarities, i.e., we assumed that 156 similarities between individuals within a population are greater than those between individ-157 uals belonging to different populations, and that parameters representative of the population 158 could be obtained. 159

First, we tested whether a single set of parameters (Marn et al. [19], an updated set in Table 2) can describe both populations. This parameter set will be henceforth referred to as $'pars_{NA}'$. We simulated environmental conditions (temperature and scaled food availability) of the Mediterranean Sea, therefore testing whether environment alone can account for the observed differences between the two populations (Fig. B.13).

Second, we assumed that some parameters might differ between the two populations, and then estimated these parameters using only data for the Mediterranean population and ' $pars_{NA}$ ' as initial values. Out of the 15 core parameters of the standard DEB model (includes the extra maturity level for hatching), we allowed eight parameters to differ between populations and used a species-specific value or a default value for the remaining parameters (see Table 2). The resulting parameter set will be henceforth referred to as ' $pars_{MED}$ '.

We quantified the goodness of fit between each model prediction and the corresponding 171 data set or data point with relative error (RE) and symmetric squared error (SSE). Calcu-172 lating the mean of all RE and the mean of all SSE gave two measurements for an overall 173 goodness of fit (MRE and SMSE, respectively) [58, this issue]. We assessed the performance 174 of the two parameter sets by comparing the corresponding MRE and SMSE values. In addi-175 tion, we compared RE values obtained by ' $pars_{NA}$ ' to RE values obtained by ' $pars_{MED}$ ' using 176 the Wilcoxon signed rank test (signrank in Matlab) and sign test, which have been rec-177 ommended for comparisons of two classifiers over multiple data sets [59]. We approximated 178 uncertainty of parameter values by combining the information on data availability (COM-179 PLETE level, [57]) and data agreement (MRE,SMSE), following the approach presented in 180 Pauly and Zeller [60]. 181

We hypothesized that if model predictions based on $'pars_{MED}'$ fit the data better than those based on $'pars_{NA}'$ after accounting for environmental differences, then we would be able to infer which physiological characteristics cause the two populations to differ. Put alternatively, we would better understand how evolution adapted MED loggerhead turtles to life in the Mediterranean Sea. We wrote and executed all code in Matlab R2015a, using the current release of the DEBtool package ([61], downloaded 09-April-2018) for parameter estimation [58, this issue] and model predictions.

Gathering data. The process of parameter estimation uses several types of data [57, 58, 62, 189 this issue]: data-*points* such as age and size at hatching and puberty, maximum reproduction 190 etc.; and data-pairs such as length-at-age, mass-at-length, etc. We obtained the necessary 191 data by performing a comprehensive literature search and securing previously unpublished 192 data from a rearing program in Marineland. The latter data pertained to adults captured 193 in the Mediterranean Sea and their offspring. Generally, we prioritized straight carapace 194 length (SCL) [63]. We did use curved carapace length (CCL) to estimate length at puberty 195 and ultimate length, because only CCL had been reported in most published literature on 196 Mediterranean loggerhead turtles. Some data required additional assumptions (stated below); 197 all data is presented in Table 3, and Figs. 3–6. 198

Birth from the perspective of physiological energetics denotes the transition from the embryo stage, in which there is no feeding nor reproduction, to the juvenile stage, in which feeding occurs. In the case of loggerhead turtles, hatching as the moment of leaving the egg, emergence as the moment of leaving the nest, and birth as the onset of feeding take place several days apart. We assumed that the duration of the period between hatching and birth depends on temperature, but is the same for both populations.

We considered sexual maturity or puberty, defined as the onset of energy allocation to 205 reproduction, as equivalent to the event of first nesting (as in Marn et al. [19]). Only estimates 206 were available for the age at puberty because the onset of energy allocation to reproduction 207 generally cannot be observed directly. Estimates are reported as a range and mostly depend 208 on the length defined as the "length at puberty" and on the method used for estimation 209 [3, 6, 7, 64]. The value used as the "observed value" (18 years) was therefore quite uncertain 210 and was consequently given lower weight in the parameter estimation procedure (see also 211 Lika et al. [57], Marques et al. [58, this issue]). We calculated size (length and wet weight) 212 at puberty as the average of the low end of the reported ranges. 213

To calculate the reproductive output (clutch size) of a female of a certain size, we modeled the investment into reproduction as a continuous process and then used a simplified linear relationship by which females produce two clutches per nesting season every two years (average values from Broderick et al. [35]). We calculated the maximum reproductive output using the high end of the reported median values (2.2 clutches per nesting season with nesting 2 years apart).

We assumed the maximum life span and the maximum length under *ad libitum* food to be

relatively consistent within a species and, due to a lack of data for MED loggerhead turtles, we used the same values as for the NA population: 65 yr [65, 66] and 130 cm SCL [67]. We calculated the ultimate size as the average of the high end of the reported values. We then used the ratio between the average ultimate length and maximum length as the initial value for scaled food availability (f = 0.67) during parameter estimation.

All of the aforementioned data-*points* described the MED loggerhead turtles living and 226 nesting in the wild. Datasets of data-*pairs* only partially pertained to the wild MED turtles: 227 temperature vs. incubation duration, growth in length, and carapace length vs. clutch size. 228 We secured additional growth data (mass and carapace length vs. time) from a rearing 229 facility located in the Mediterranean. Data from the rearing facility comprised four years of 230 measurements of two loggerhead turtles that hatched in 2010, and three years of measurement 231 of several loggerhead turtles that hatched in 2011. An important feature of these data was 232 that the exact age of measured individuals and environmental conditions in terms of food 233 and temperature had been known. 234

Following the approach for the NA population [19], we characterized the environmental 235 conditions experienced by wild loggerhead turtles by an average sea surface temperature and 236 average food availability. Specifically, we set $T_{\rm MED} = 21^{\circ} C$ [24] because this is the average 237 seawater temperature in the eastern Mediterranean basin where MED loggerhead turtles are 238 mostly concentrated [1]. For food availability, we initially set f = 0.67 based on the ratio 239 between ultimate and maximum lengths, but later we treated this quantity as a parameter to 240 be estimated simultaneously with other parameters. We characterized the rearing facilities 241 with recorded temperature $(T = 22 - 26^{\circ}C)$ and *ad libitum* food (f = 1). Because we could 242 include into the model different temperatures and food availabilities for wild and captive 243 individuals, it was possible to simultaneously use data from the wild and from captivity. 244

245 3. Results

246 3.1. Size and condition of Mediterranean and North Atlantic loggerhead turtles

Statistical analyses point to a distinct physiology between the two studied populations 247 (Fig. 1, Table 1). On average, MED eggs are 14% shorter and 45% lighter than NA eggs, MED 248 hatchlings are 11% shorter and 23% lighter than NA hatchlings, and nesting MED females are 249 19% shorter and 64% lighter than nesting NA females. Surprisingly, despite these differences 250 in size and mass between the two populations, condition remains similar when comparing the 251 same life events (Table 1). Such similarity, taken together with the decrease of the condition 252 index with age (Table 1), suggests that condition may be a geographically invariant indicator 253 of the ontogenetic state of loggerhead turtles. We could only obtain point estimates for the 254

²⁵⁵ condition indices in Table 1, which is why we refrained from using them in any additional ²⁵⁶ statistical analyses or to make any definite conclusions. However, we did have independent ²⁵⁷ data to estimate the 95% confidence intervals for the condition indices of NA hatchlings and ²⁵⁸ adults (Fig. 2). The obtained values are in broad agreement with the results in Table 1.

Table 1: Population-level means (and standard deviations) of egg diameter and wet weight, SCL and wet weight at hatching, and SCL and wet weight at nesting. Fulton's condition index, K, is given for comparison (Eq. (1)). The condition index is similar if the same life events of the two populations are compared, but seems to decrease through ontogeny. See Methods and Tables A.4-A.6 in Appendix A for more details.

Life stage	Measurement	North Atlantic	Mediterranean
Egg size	Diameter (cm)	$4.25 (SD \ 0.14)$	$3.71 (SD \ 0.224)$
	Wet weight (g)	$41.98 (SD \ 3.00)$	$28.90 (SD \ 3.241)$
	K	0.5469	0.5659
Hatching	Length SCL (cm)	$4.51 (SD \ 0.1456)$	4.05 (SD 0.376)
	Wet weight (g)	$19.26 (SD \ 1.642)$	$15.60 (SD \ 1.933)$
	K	0.2100	0.2348
Nesting	Length SCL (cm)	$92.84 (SD \ 4.539)$	78.03 (SD 5.138)
			$78.77 \text{ (SD } 4.521)^{\dagger}$
	Wet weight (kg)	$110.18 (SD \ 18.054)$	$67.26 \text{ (SD } 8.62)^{\dagger}$
	K	0.1377	0.1376†

Condition of two adult females in the Marineland (France) reproductive program is K = 0.1959 and K = 0.2525.

[†] Values based on data pertaining to the turtles nesting in Greece and Cyprus only, which are slightly larger than those nesting in, e.g., Turkey and Libya. Marked values were used to calculate the condition index of nesting Mediterranean loggerhead turtles, because Wet weight data was available just from Greece and Cyprus.

Further statistical analyses are outlined in Appendix A (Figs. A.9–A.11) with the main result being a larger size of NA turtles relative to MED turtles almost irrespective of the compared locations. However, given the limitations in data quality and a lack of mechanistic underpinning for the obtained results, we proceed to examine the observed size differences between MED and NA loggerhead turtles by means of a modeling approach.

264 3.2. Mechanistic DEB-based modeling of the Mediterranean population

Our approach yielded two sets of model predictions that we compared to the data. First were the predictions obtained by assuming that the same parameter values pertained to both NA and MED populations (' $pars_{NA}$ ' in Table 2 excluding the value of scaled functional response, f), meaning that the differences between the two populations originated solely from the different environmental conditions. Second were the predictions obtained by estimating the parameter values for MED population (' $pars_{MED}$ ' in Table 2), and then accounting for the different environmental conditions. The relative errors for predictions obtained with

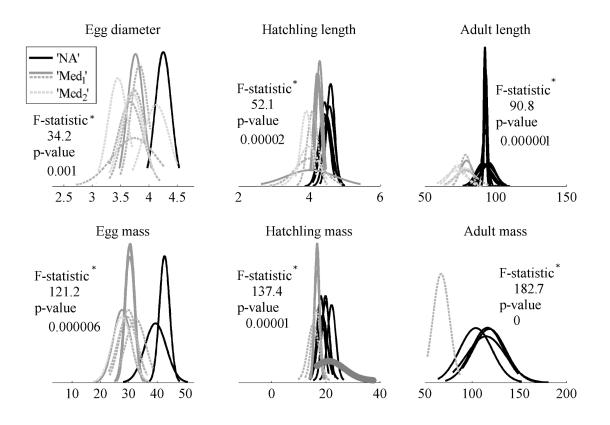


Figure 1: Theorized size distributions of loggerhead turtles and the results of the Welch ANOVA test. The ANOVA in all cases rejects the hypothesis that all samples come from the same distribution. A visual inspection of size distributions suggests that NA turtles are larger than MED ('Med₁'-Greece and Cyprus; 'Med₂'-Turkey and Libya) irrespective of the life stage. Normal size distributions were assumed and plotted using means and standard deviations from Tables A.4–A.6. Where a value for the standard deviation was unavailable, we made an estimate following Hozo et al. [51]. Thick gray curve in the panel for hatchling mass represents the data from the reproduction program, which were analyzed separately. Distributions were plotted either over the reported data range (when available), or over a range defined as three standard deviations from the mean. For more details, see Figures A.9 and A.10 in Appendix A.

²⁷² 'pars_{NA}'and 'pars_{MED}' differ significantly (Wilcoxon sign rank test, F = 166, z-value = -2, ²⁷³ p=0.004). Predictions with the 'pars_{MED}' parameter set perform better in terms of the sign ²⁷⁴ test, wherein 19 (57.58%) data sets have a significantly smaller relative error with 'pars_{MED}' ²⁷⁵ than 'pars_{NA}'. The 'pars_{MED}' parameter set also yields smaller MRE and SMSE values of ²⁷⁶ 0.138 and 0.222 relative to 0.194 and 0.349 for 'pars_{NA}'. Keeping in mind that the life history ²⁷⁷ traits observed in nature are better defined in terms of ranges than unique critical values, we ²⁷⁸ focus on qualitative rather than numerical analysis of the differences between populations.

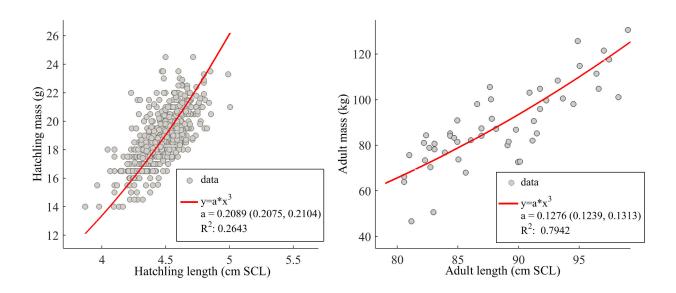


Figure 2: Estimating the condition indices of NA loggerhead turtles with independent data. By fitting model $y = ax^3$, where x-axis is length, and y-axis is mass, parameter a becomes an estimate of the condition index (95% confidence intervals in parentheses). The R^2 is satisfactory for the adult data, but low for the hatching data. The low R^2 could signify a large inter-individual variation, but could also be an artifact of the measurement error as the ranges of measured weights and lengths are small (10 g and 1 cm, respectively). Data sources: size at hatching partially published in Stokes et al. [37]; size at nesting from Wabnitz and Pauly [68] selected such that length is larger than the smallest reported length of nesting turtles [69, 70].

Table 2: Parameters for the North Atlantic (' $pars_{NA}$ ', AmP *Caretta caretta* [dataset] 2018) and the Mediterranean (' $pars_{MED}$ ') loggerhead turtle at reference temperature $T_{ref} = 273$ K. Primary parameters (directly linked to processes), auxiliary parameters (linking measured and observed quantities), and compound parameters (functions of two or more primary parameters, [16]) are separated with horizontal lines. Parameters estimated using the covariation method [57, 58, this issue] are marked with the † symbol. Parameters marked with a **bold** font are later discussed in more detail. Primary and auxiliary parameters for which the default values were used are listed below the table. More details about parameter values of loggerhead turtles can be found in Marn et al. [19], N/A [20].

Parameter	Symbol	pars_{NA}	' $\operatorname{pars}_{\operatorname{MED}}$ '	Unit
Maximum specific assimilation rate	$\{\dot{p}_{Am}\}$	747.3^{\dagger}	895^\dagger	$\mathrm{Jd^{-1}cm^{-2}}$
Energy conductance	\dot{v}	0.0681^{\dagger}	0.0721^{\dagger}	${\rm cm}{\rm d}^{-1}$
Allocation fraction to soma	κ	0.729^{\dagger}	0.733^{\dagger}	-
Volume-specific somatic mainte- nance	$[\dot{p}_M]$	11.20^\dagger	13.5^\dagger	$\mathrm{Jd^{-1}cm^{-3}}$
Specific cost for structure ^a	$[E_G]$	7322	7322	$\rm Jcm^{-3}$
Maturity maintenance rate coefficient	\dot{k}_J	0.00112^{\dagger}	0.00152^{\dagger}	d^{-1}
Maturity at hatching ^b	E_H^h	21080^\dagger	22320^\dagger	J
Maturity at birth ^b	$E_H^{\overline{b}}$	25350^\dagger	22320^\dagger	J
Maturity at puberty	E_H^p	$old 98750^\dagger$	58580^\dagger	kJ
Weibull aging acceleration	\dot{h}_a	$1.112\text{e-}10^\dagger$	$1.532\mathrm{e}\text{-}10^\dagger$	d^{-2}
Scaled functional response	f	0.81^{\dagger}	0.67^{\dagger}	_
Arrhenius temperature ^a	T_A	7200	7200	Κ
Shape coefficient	$\delta_{ m SCL}$	0.3915^{\dagger}	0.3915	-
Shape coefficient ^c	$\delta_{ m CCL}$	$0.3117^{\dagger} \ (0.3645)$	0.3645^{\dagger}	-
Density of structure and reserve ^a	$d_V = d_E$	0.28	0.28	-
Zoom factor ^d	z	48.61^{\dagger}	48.61^{\dagger}	-
Maximum reserve density	$[E_m]$	10 980	12 410	$\rm Jcm^{-3}$
Contribution of reserve to biomass	ω	1.70	1.93	-
Maintenance ratio ^e	k	0.73	0.83	-

Other primary parameters (default values from N/A [20] used): Maximum searching rate, $\{\dot{F}_m\} = 6.51 \,\mathrm{d}^{-1} \,\mathrm{cm}^{-2}$; Digestion efficiency (of food to reserve), $\kappa_X = 0.8$; Defaecation efficiency (of food to faeces), $\kappa_P = 0.1$; Reproduction efficiency, $\kappa_R = 0.95$; Gompertz stress coefficient, $s_G = 0.0001$, Surface-area specific somatic maintenance, $\{p_T\} = 0 \,\mathrm{J/d} \,\mathrm{cm}^{-2}$.

^a Species-specific values: $[E_G]$ such that it results in 80% growth efficiency [16, 20], d_V from Kraemer and Bennett [72]; T_A estimated independently from data [1, 16, 37].

^b A delay in onset of embryonic growth (t_0 , [73, this issue]) was estimated as 12.79 d for NA and 8.89 d for MED population at $T_{ref} = 273$ K.

^c CCL shape coefficients convert different types of carapace length: for the MED population the curved carapace length has been reported, whereas for the NA population the type of length measurement has not been reported. For the results presented here, we used $\delta_{CCL} = 0.3645$ in all model runs.

^d We constrained parameter z of the MED population to the same or smaller values than that of NA population to reflect that MED individuals do not become larger than NA individuals.

^e $k = \dot{k}_J [E_G]/[\dot{p}_M]$; values larger than one are unrealistic because they imply individuals undergo metabolic switches at larger sizes when food is restricted. The only two sea turtles species in the AmP collection with values of \dot{k}_J at the default value (AmP *Dermochelys coriacea* [dataset] 2015 and AmP *Lepidochelys kempii* [dataset] 2015) have a value of k around 0.75. Predictions using ' $pars_{NA}$ ' parameter set. Parameter set ' $pars_{NA}$ ' obtained originally for NA loggerhead turtles does not match the data on MED loggerhead turtles well (3rd column of Table 3 and dashed lines in Fig. 3–6). The most important mismatches between data and predictions are (i) the failure to capture observed smaller sizes at puberty of MED individuals, and (ii) a very low prediction of the maximum reproductive output (Table 3, Fig. 6.b). The model also predicts that captive reared individuals grow slower than observed (Fig. 4).

Table 3: Comparison between observations and model predictions using two sets of parameters. Here, we set $T = 21^{\circ}$ C [24] and f = 0.67 in conjunction with both parameter sets, which are the environmental conditions prevailing in the Mediterranean. Overall goodness of fit is expressed as mean relative error (MRE) and symmetrical mean squared error (SMSE). All parameters are listed in Table 2. For more details see Section 2.2.

Data	Ob- served	$\begin{array}{l} {\rm Predicted} \\ {\rm ('pars_{NA}')} \end{array}$	$\begin{array}{c} \text{Predicted} \\ \text{('pars}_{\text{MED}}\text{'}) \end{array}$	Observed, range	Unit	Data source
age at hatching	49.08	51.06	49.58	45.8-55.8	d	1
age at birth	55.18	66.13	58.07	2-3 d after emergence	d	2,3
SCL at birth	4.04	5.36	5.17	2.5 - 4.9	cm	1, 4
wet weight at birth	15.59	19.59	18.97	9.4 - 21.5	g	4
age at puberty	18	21.7	12.41	14-28	yrs	$5,\!6$
SCL at puberty	62.5	77.07	67.8	55-69	cm	4,7,8
CCL at puberty	69.00	82.46	72.8	60-78	cm	4
wet weight at puberty	52.00	58.07	42.71	52.5	kg	7
life span	67.00	75.5	67	65+	yrs	9
ultimate SCL	87.00	83.32	82.85	77-91	cm	4,7,8
ultimate CCL	91.00	89.14	88.96	85-99	cm	4
ultimate wet weight	87.00	73.35	77.79	87	kg	7
initial energy content of the egg	170.00	164.5	156.6	165-260	kJ	10
maximum reproduction rate †	0.4822	0.2022	0.5158	0.3452- 0.8630	d^{-1}	4,7,8,11
MRE	-	0.194	0.138	$[0 \infty)$		
SMSE	-	0.349	0.222	$[0 \ 1]$		

[†] Expressed as the number of eggs per day by taking into account the 2.2 nests (clutches) per nesting season, 160 eggs per clutch, and remigration interval of 2 years: $R_i = 2.2 \times 160/(2 \times 365) d^{-1}$. **Data sources**: {1} Reid et al. [38], {2} Godfrey and Mrosovsky [76], {3} data partially published in Stokes et al. [37], {4} Margaritoulis et al. [1], {5} Casale et al. [7], {6} Piovano et al. [3], {7} Groombridge [11], {8} Tiwari and Bjorndal [5], {9} FAQ https://gstc.jekyllisland.com/, Accessed: 18-Nov-2015, {10} Hays and Speakman [8], {11} [35]

Predictions using ' $pars_{MED}$ ' parameter set. The parameter set estimated for MED loggerhead turtles (Table 2) generates close-to-the-observed size at puberty, reasonable reproductive output, and fast growth of captive reared individuals at different temperatures (4th column of Table 3 and solid lines in Figs. 3–6). Most of the predictions for life history traits are within or close to values reported for MED loggerhead turtles (Table 3).

Furthermore, age at hatching and birth is predicted well (Table 3), as is the slope of incubation duration as a function of temperature, although the accompanying intercept is somewhat underpredicted (Fig. 3). The predicted SCL and wet weight at birth are larger than observed, whereas the predicted wet weight and age at puberty are lower than observed (Table 3).

The model also matches growth and the length-to-weight relationship for captive-reared juveniles, and faster growth at higher temperatures (Figs. 4 and 5). In reality, very similar conditions sometimes lead to markedly different growth patterns (e.g., see Fig. 4, top right and bottom left panels), which contributes substantially to the overall mean relative error.

Modeled growth of wild MED turtles matches the data up to approximately 25 cm CCL and after 70 cm CCL, but there is a mis-match in-between (Fig. 6, left panel). Clutch size as a function of carapace length (Fig. 6) suggests a more complex relationship than the assumed two clutches every two years. The predictions for clutch size are realistic (90-160 eggs per clutch), but initial energy in an egg is a bit lower than Hays and Speakman [8] assumed for MED loggerhead turtles (Table 3).

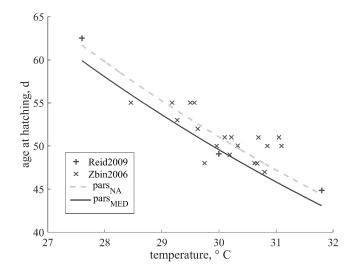


Figure 3: Incubation duration as a function of incubation temperature of wild MED loggerhead turtles - data and model predictions. Legend denotes the parameter set used to obtain model predictions (Table 2, see text for more details). The scaled functional response was estimated to be 0.703 for the MED population. Data from Reid et al. [38].

Model parameters. All nine estimated primary parameters $(\{\dot{p}_{Am}\},\kappa, [\dot{p}_{M}], \dot{v}, E_{H}^{h}, E_{H}^{b}, E_{H}^{b}, k_{J}, \text{ and } h_{a})$ differ in value between populations (Table 2). With a relatively high level of

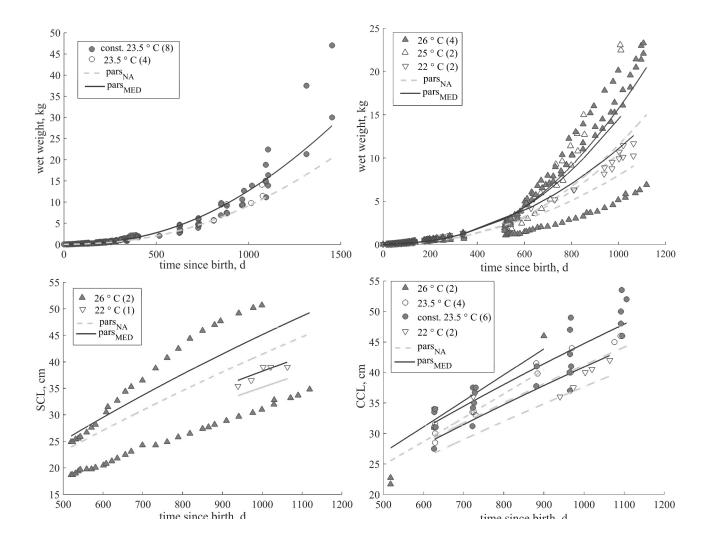


Figure 4: Growth of captive reared hatchlings. The top two panels show the predicted growth in wet weight, while the bottom two panels show the predicted increase in carapace length with age. Different symbols indicate different temperatures at which data was collected, and numbers in brackets refer to the number of individuals kept at a specific temperature regime (see legend). Parameter set used to obtain a specific set of model predictions (excluding f) is denoted in the legend (values in Table 2, see text for more details). Temperature was explicitly included in model simulations and resulted in steeper growth curves predicted for higher temperatures. Food was assumed *ad libitum* (f = 1). Previously unpublished data from Marineland rearing facility.

completeness ≈ 3 and a satisfying goodness of fit in the context of the AmP collection [21], we judge the level of uncertainty of parameter values to be $\approx 10-20\%$ (Table 2 in Pauly and Zeller [60]). Differences exceeding this threshold are present in three primary parameters: specific assimilation rate ($\{\dot{p}_{Am}\}$), specific somatic maintenance rate ($[\dot{p}_M]$), and maturity level at puberty (E_H^p) (Table 2).

Specific assimilation rate $(\{\dot{p}_{Am}\})$ and specific somatic maintenance rate $([\dot{p}_M])$ are 20% higher for MED loggerhead turtles than for their NA relatives. The surface-area specific

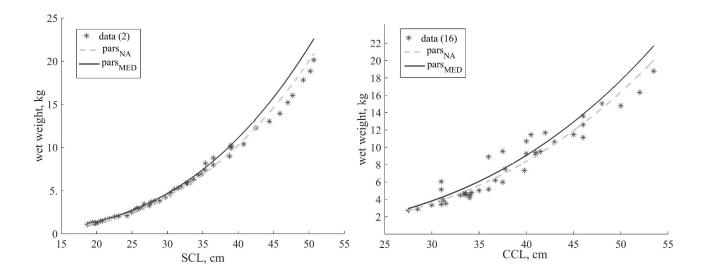


Figure 5: Wet weight as a function of length for captive reared hatchlings. Numbers in brackets refer to the number of individuals in the dataset (see legend); predictions are independent of temperature. Parameter set used to obtain a specific set of model predictions (excluding f) is denoted in the legend (values in Table 2). Food was assumed *ad libitum* (f = 1). Previously unpublished data from Marineland rearing facility.

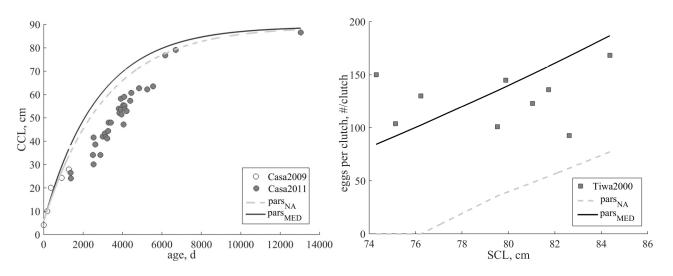


Figure 6: Growth in length (left panel, data from Casale et al. [7, 77]), and clutch size as a function of carapace length (right panel, data from [5]) of wild MED loggerhead turtles: data and model predictions. Growth is overpredicted between sizes of approximately 25 and 65 cm CCL. The linear relationship between carapace length and clutch size (assuming an average number of 2 clutches per nesting season every 2 years) does also not completely match the data. Predictions were obtained for the MED environment assuming the (constant) estimated f_{MED} (Table 2) and $T = 21^{\circ}$ C; mis-match with the data suggests a more variable environment and/or additional factors which were not included in the calculations. Parameter are listed in Table 2; see text for more details.

assimilation rate is a primary parameter calculated as $L_m^{ref} z[\dot{p}_M]/\kappa$, where L_m^{ref} is a reference

structural length of 1 cm. The zoom factor, z, is tightly linked to the maximum length an 316 animal can reach. Since NA individuals are typically larger than MED individuals [6], the 317 zoom factor z was estimated with the constraint that it cannot be larger for the MED than 318 the NA population. When compared to the ' $pars_{NA}$ ' set, z is the same and allocation to soma 319 κ is similar, but somatic maintenance rate $[\dot{p}_M]$ is higher in the 'pars_{MED}' set. Consequently, 320 the parameters imply that MED loggerhead turtles can assimilate 20% more energy per unit 321 of structural surface area per day than NA loggerheads can $(\{\dot{p}_{Am}\}=895 \,\mathrm{J \, cm^{-2} \, d^{-1} \, compared})$ 322 to $747 \,\mathrm{J}\,\mathrm{cm}^{-2}\,\mathrm{d}^{-1}$), but need to pay 20% more maintenance per unit of structural volume per 323 day $([\dot{p}_M] = 13.5 \,\mathrm{J \, d^{-1} \, cm^3}$ compared to $11.20 \,\mathrm{J \, d^{-1} \, cm^3}$). 324

Maturity levels (hatching E_H^h , birth E_H^b , and puberty E_H^p) all have lower values for the MED population, with the biggest difference of 40% for E_H^p (58 580 kJ in '*pars*_{MED}' compared to 98 750 kJ in '*pars*_{NA}'). Maintaining each unit of maturity is slightly more energy-expensive for MED turtles than for NA turtles (k_J =0.0015 J d⁻¹ compared to 0.0011 J d⁻¹).

The three primary parameters with inter-population differences exceeding the 20% thresh-329 old are linked to three processes: assimilation, maintenance, and maturation (reproduction). 330 Energy budget of MED turtles changes through ontogeny, and so does the daily energy allo-331 cation between main physiological processes (Fig. 7). The energy allocation exhibits a similar 332 pattern between the two populations, with the similar proportions of daily assimilation allo-333 cated to maintenance and development (in juveniles), or reproduction (in adults) (numbers 334 above the bars in the lower panel of Fig. 7). However, the absolute amounts of energy assimi-335 lated and invested daily into specific processes differ. For example, because MED loggerheads 336 grow to a smaller size, a fully grown MED female daily needs to invest $\approx 270 \, \text{kJ}$ less energy 337 into somatic and maturity maintenance than a fully grown NA female. Even though the 338 MED female also assimilates less energy ($\approx 340 \, \text{kJ.d}^{-1}$ less than the NA female), she can still 339 invest 12% of her assimilation flux into reproduction, which is comparable to 16% invested 340 by the NA female (lower panel in Fig. 7). 341

MED loggerhead turtles have slightly larger energy conductance (\dot{v} =0.0721 cm d⁻¹ in 342 'pars_{MED}' compared to 0.0681 kJ in 'pars_{NA}'). Their maximum reserve density ($[E_m]$ = 343 $\{\dot{p}_{Am}\}/\dot{v}\}$ is however larger (12.4 kJ cm⁻³ in 'pars_{MED}' compared to 11 kJ cm⁻³ in 'pars_{NA}'), 344 resulting in a larger contribution of reserve to biomass ($\omega = 1.93$ in 'pars_{MED}' compared to 1.70 345 in 'pars_{NA}', Table 2). These results are interesting in the context of similar condition indices 346 (Table 1) regardless of the lower food availability in the Mediterranean Sea (see Discussion). 347 Initially, based on the ratio between maximum and ultimate lengths, we set scaled food 348 availability to f = 0.67. By modeling all life history points (Table 3) and multiple datasets, 349 such as reproduction and growth (Fig. 6) of wild individuals we found that f = 0.67 represents 350 the feeding conditions in the Mediterranean very well. However, because similar condition 351

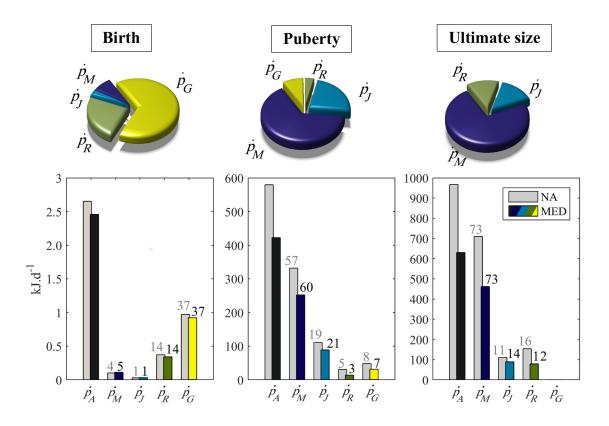


Figure 7: A visualization of the MED loggerhead turtle's energy budget at birth, puberty, and ultimate size: \dot{p}_A - assimilation flux, \dot{p}_G - growth flux, \dot{p}_R - maturation/reproduction flux, \dot{p}_M - somatic maintenance, and \dot{p}_J - maturity maintenance, presented as fractions of the mobilization flux (pie charts) and as absolute values in kilojoules (histograms). Numbers above the histogram bars are proportions of the assimilation flux invested into a specific process, with NA values denoted in gray and MED values in black. Values do not necessarily add to a 100% - energy that is assimilated but not mobilized stays in the reserve. Legend: MED - Mediterranean loggerhead turtle (parameters $pars_{\text{MED}}, f = 0.67$), NA - North Atlantic loggerhead turtle (parameters $pars_{\text{NA}}, f = 0.81$). Parameters values are listed in Table 2.

indices of NA and MED individuals (Table 1) hint at a similar scaled food availability, we repeated the parameter estimation starting with the value of f = 0.81 (from 'pars_{NA}') to ensure robustness with respect to the starting choice of f. The estimation routine, which could now freely vary the value of f, returned f = 0.67, thus confirming the validity of our original setting.

357 4. Discussion

Environmental conditions can affect physiological characteristics of organisms. For example, reducing food availability directly reduces ultimate size, and changes in temperature directly affect the growth rate [12, 14, 78]. The environmental effects can also be indirect. Even subtle differences in environments can affect individuals on the biomolecular level and affect energy utilization patterns. We have found evidence of both direct and indirect effectsin this study.

We analyzed loggerhead turtles from the Mediterranean (MED) population, and compared them to loggerhead turtles from the North Atlantic (NA) population. By simultaneously analyzing morphology, physiological (condition) index, and the energy budget-related processes of turtles inhabiting areas with different environmental conditions, we explored how environmental conditions, physiological characteristics, and the interaction of the two might result in size dimorphism and different growth and reproduction reported previously [3, 5, 6].

370 4.1. Analyses of size and condition indices

Empirical analyses show that the size dimorphism is present during the whole loggerhead 371 turtle life cycle, with MED turtles being smaller than NA loggerhead turtles. Captive-reared 372 hatchlings are heavier than the wild hatchlings of both populations (Figure A.9, panel b), 373 which is consistent with the maternal effect as implemented in the standard DEB model; 374 better fed mothers produce larger offspring [73, 79, this issue]. Size dimorphism is most 375 pronounced at nesting; size difference between nesting females is consistent with a previous 376 study [5, 6]. Because food availability is lower in the Mediterranean Sea than in the North 377 Atlantic, [23, 80, 81], the dimorphism could qualitatively be explained by this difference 378 [14, 16]. But is it enough? 379

The differences in food availability may not be the only reason for size dimorphism. We 380 would intuitively expect an organism exposed to less food in the environment to have a lower 381 condition index. The two populations, however, have similar condition indices regardless of 382 the lower food availability in the Mediterranean Sea. To see whether our model provides 383 an explanation for this pattern, we refer back to Eqs. (2) and (3) to obtain a first-order 384 theoretical approximation for the condition index, $K = \delta^3 \rho (1 + f\omega)$. Assuming the same 385 shape factor, δ , and density of structure, ρ , for turtles from both populations (as we did 386 throughout this study) shows that the ratio of condition indices is 387

$$\frac{K_{\rm NA}}{K_{\rm MED}} = \frac{1 + \omega_{\rm NA} f_{\rm NA}}{1 + \omega_{\rm MED} f_{\rm MED}}.$$
(4)

³⁸⁸ NA loggerhead turtles experience better feeding conditions, $f_{\rm NA} = 0.81$ compared to $f_{\rm MED} =$ ³⁸⁹ 0.67. In contrast, MED loggerhead turtles have a higher maximum reserve density, leading ³⁹⁰ to $\omega_{\rm NA} = 1.70$ compared to $\omega_{\rm MED} = 1.93$ (Table 2). Inserting these values into Eq. (4) gives ³⁹¹ $K_{\rm NA}/K_{\rm MED} = 1.0366$, thus indicating that the conditions indices of the two populations should ³⁹² be very similar. More importantly, we see that the reserve capacity of MED loggerhead turtles ³⁹³ compensates for lower food availability in the Mediterranean ([E_m], Table 2). Increasing the reserve capacity is, in fact, a reasonable adaptation to lower food availability because organisms with more reserves better withstand starvation periods, which in turn are more likely to occur in an environment with little food to begin with. Admittedly, Eq. (4) misses potentially important contributions to the condition index, such as the reproductive buffer or the total body water content, but these contributions are variable in time, and thus responsible for second-order effects for which more elaborate longitudinal measurements are necessary.

401 4.2. Environmental characteristics known to affect size and physiological performance

Incubating (nest) environment can be very different between NA and MED nesting 402 beaches, and may be responsible for some of the size difference in hatchlings. Based on 403 results from intra-population studies, incubation at colder and moister sites generally yields 404 heavier and larger hatchlings [37, 40, 41, 42, 43], but see [38, 42, 45, 46]. In contrast, high 405 salinity in the incubating environment, mirroring the physiological effects of a dry incubat-406 ing environment, yields smaller hatchlings [44]. Nesting locations for each of the populations 407 show great variability in the average incubation duration, hatchling size, and hatchling sex 408 ratio [82], suggesting markedly different incubation conditions even within a single popu-409 lation. Making comparisons on the basis of population-specific abiotic factors is therefore 410 extremely susceptible to source-based bias in the available data. With caution, nevertheless, 411 some general patterns can be explored. 412

The average incubation temperatures of nests on MED beaches (Cyprus and Turkey, [83, 413 84]) are higher than those on NA beaches (Georgia, [85]), the higher temperature in natural 414 nests being often combined with a drier substrate [42]. Such a combination suggests that the 415 smaller size of the MED hatchlings could be linked to warmer and drier nests. The correlation 416 of temperature and humidity to size at hatching does, however, exhibit some inconsistencies 417 (cf. Stokes et al. [37], Reece et al. [42], Glen et al. [43], Ji and Brana [45], Patino-Martinez 418 et al. [46]). In addition to temperature and moisture affecting the hatchling morphology 419 independently, a combination of the effects can be present [40], including indirect effects on 420 size via incubation duration and embryonic volk utilization [40, 45, 86]. 421

Osmotic water absorption of eggs and embryos of freshwater snapping turtle is susceptible to salinity [44]. It is reasonable to assume that osmotic absorption of sea turtle eggs and embryos [46] is susceptible to similar phenomena. However, the effects of salinity on embryonic development and hatchling size have yet to be explored for loggerhead turtles.

Higher food availability and warmer temperature should both positively affect growth, reproduction, and size of loggerhead turtles [14]. We assumed the same temperature sensitivity within the thermal tolerance range (parameter T_A) for individuals from both popula-

tions. The assumption was supported by the good fit of model predictions to data (Figs. 3) 429 and 4). Food availability was assumed higher in the North Atlantic. Within the same popu-430 lation, females that experience higher food availability are larger, reproduce more frequently 431 [14, 87, 88], and produce larger hatchlings [88]. However, the egg size within a population is 432 generally uniform [5] and does not differ significantly in (energy-providing) yolk content [89]. 433 Eggs of different populations do, however, differ in size with a positive correlation between 434 volk amount and egg size (e.g., [50]). Earlier onset of reproduction is linked to food availabil-435 ity in sibling species [90]. It therefore seems reasonable to assume that such environmental 436 cues may cause distinct adaptations at the population level. 437

Other pressures present in the environment, such as predation and anthropogenic pres-438 sures, can result in evolutionary size selection. Smaller size and younger age at nesting, as 439 identified in this study (see also Piovano et al. [3]), might be a result of the selection driven by 440 anthropogenic pressures. Strong anthropogenic pressure (especially, commercial harvesting) 441 is known to drive down the age and size at first breeding of fish and predators in the marine 442 ecosystems [91, 92]. The main source of loggerhead turtle mortality in the Mediterranean is 443 by catch [11, 93], with other significant anthropogenic pressures such as shipping and mass 444 tourism [94] also having a high probability of interaction and a negative impact on loggerhead 445 turtles. Such pressures exert a strong evolutionary incentive for the MED population to breed 446 at a younger age and, consequently, smaller size—particularly in a food-limited environment. 447

448 4.3. Model predictions for MED loggerhead turtles

Given the relatively high level of data completeness [21], our model generates satisfactory fits, but it is still imperfect. The mis-matches between the model's outputs and data are informative nonetheless, because they help identify limitations in data availability or oversimplifications used in the model formulation.

For example, the "observed" size at nesting, calculated from all available data, included a relatively high proportion of sea turtles nesting in Turkey and Libya, which are smaller than those nesting in Greece and Cyprus (Margaritoulis et al. [1], Groombridge [11] and Table A.6). In reality, the Greek subpopulation comprises the largest proportion of the Mediterranean population [1], meaning that the average length at puberty is likely to be larger, i.e., closer to the predicted value.

Age at puberty was predicted to be lower than the observed value, as in other sea turtle DEB models [19, 73, this issue]. It could be that sea turtles start allocating to reproduction several years prior to actual nesting, or that they experience a period of less favorable (than simulated) environmental conditions during their juvenile stage, which would prolong the time needed to reach puberty [14, 95]. A less favorable environment during the juvenile stage would account also for slower-than-predicted growth of wild MED turtles in Fig. 6 (leftpanel).

A favorable prediction of the maximum reproduction rate (Table 3) and a somewhat less satisfactory prediction of the clutch size as a function of carapace length (Figure 6) imply that the modeled allocation to reproduction is correct, but the clutch size is determined by factors other than available energy alone. Data for total seasonal reproductive output rather than clutch size may be more appropriate for model validation, but are not available in part because of how logistics hinders data gathering [5, 9, 10].

In summary, our model successfully reproduces the main life-history traits and the size dimorphism of MED and NA loggerhead turtles, but only after calibrating both the environmental forcing and the model parameters. The need to separately calibrate the model parameters suggests that MED loggerhead turtles underwent evolutionary adaptations to the prevailing conditions in the Mediterranean Sea. Furthermore, the success of our modeling approach provides support for a wider implementation of predictive models in conservation efforts [96].

479 4.4. Can a North Atlantic turtle thrive in the Mediterranean Sea?

⁴⁸⁰ Applying the same theoretical and practical framework as was done for the NA population ⁴⁸¹ [14, 19], we simulated the Mediterranean environment for a NA turtle, and the North Atlantic ⁴⁸² environment for a MED turtle. Simulated environments effectively differed only in the scaled ⁴⁸³ food availability ($f_{\text{MED}} = 0.67$ this study, $f_{\text{NA}} = 0.81$ Marn et al. [19], AmP *Caretta caretta* ⁴⁸⁴ [dataset] 2018). The turtles physiology differed due to parameter values (Table 2).

Predictions for the NA loggerhead turtle imply that this turtle would reach a smaller 485 ultimate size than she would in the North Atlantic (83 cm compared to 105 cm SCL), i.e., the 486 same ultimate size as is predicted for the MED turtle in the Mediterranean. Size at puberty 487 is predicted to be similar as in the North Atlantic (77 cm SCL), however the needed time to 488 reach puberty would be 9 years longer than in the North Atlantic (22 compared to 13 years) 489 and the reproduction rate would be only about 26% of that observed in the North Atlantic 490 (Marn et al. [19] and AmP Caretta caretta [dataset] 2018). The combination of much delayed 491 onset of reproduction and a lower reproduction rate, would drastically reduce fitness of NA 492 turtles. Assuming a lower sea temperature based on the information that the NA loggerhead 493 turtles mostly forage in the western (cooler, 19°C Manca et al. [24]) Mediterranean basin 494 [2, 3] would amplify the effect further. Hence, it is possible that, even without fidelity to 495 their nesting beaches, the NA turtles visiting the Mediterranean Sea would not reproduce 496 there due to energy limitation. 497

⁴⁹⁸ The Mediterranean turtle venturing into the Atlantic would have an advantage over the

NA loggerhead turtles. Because she needs to invest less energy into maturation to reach 499 puberty than the NA turtle does (Table 2), she would mature several years earlier (pre-500 dicted age at puberty at f = 0.81 is 9 years for the MED loggerhead turtle). Lower energy 501 needed for maintaining the maturity level of puberty enables that more energy is allocated 502 to reproduction: predicted ultimate size of 100 cm SCL matches the prediction for the NA 503 turtle in the North Atlantic, and the reproduction rate at the ultimate size was predicted to 504 be around 65% higher than the observed reproduction rate of NA individuals of the same 505 size, and 43% higher than the reproduction rate predicted by the model for the NA turtle 506 (Marn et al. [19], and AmP Caretta caretta [dataset] 2018). Increase in size is also predicted 507 to be faster for the MED loggerhead turtle. Faster maturation and growth of MED turtles 508 experiencing similar condition as NA turtles is in accordance with reports [3], however there 509 are no available data on MED loggerhead turtles nesting in the North Atlantic that we could 510 use to check whether such high reproduction output is reasonable. An advantage over NA 511 turtles would be a strong incentive for MED turtles to go into North Atlantic. Perhaps they 512 do (but encounters with them have not been frequent) or the fidelity to nesting beaches and 513 direction of sea currents keep them in the Mediterranean Sea. 514

515 4.5. Evolutionary implications of the parameter values

Our results are consistent with the idea that MED loggerhead turtles have evolved from 516 NA ones. Namely, known genetic differences between the two populations [2, 3] are likely 517 to be reflected in the different parameter values in the model, because genotype determines 518 phenotype, and parameter values are a manifestation of the latter [97]. The simulations 519 presented herein were indeed unable to reproduce the different life-history traits of MED 520 and NA loggerhead turtles simply by modifying the environmental forcing of the model. 521 To successfully model both populations it was necessary to modify several core parameter 522 values (Tables 2). However, that the two populations are closely related is also seen from 523 these parameter values because they differ much less between themselves than in interspecies 524 comparisons (Fig. 8). 525

Changes in the parameter values obtained for MED loggerhead turtles relative to NA 526 ones appear consistent with the prevailing environmental conditions in the Mediterranean 527 Sea. For example, the surface area specific maximum assimilation rate $(\{\dot{p}_{Am}\})$ controls 528 the assimilation flux, but also affects energy storage capacity. A 20% higher $\{\dot{p}_{Am}\}$ for 529 MED loggerhead turtles, coupled with only a 6% higher value of the energy conductance (v, v)530 Table 2), not only enables that more energy is assimilated per unit of surface area, but also 531 results in maximum reserve density ($[E_m] = \{\dot{p}_{Am}\}/v$) of MED turtles being larger by 13%. 532 This enables MED loggerhead turtles to survive longer periods of food shortage, as expected 533

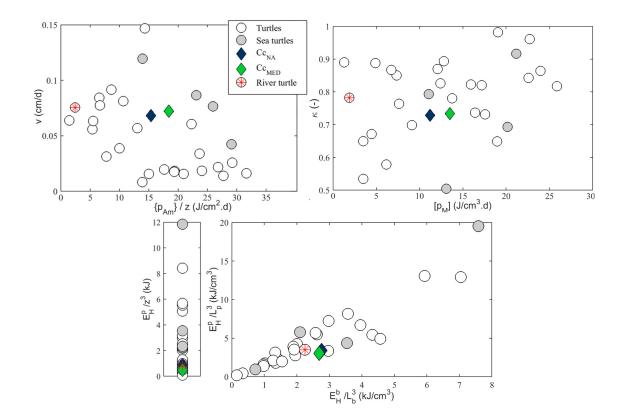


Figure 8: Values of several core parameters (maximum assimilation rate, $\{\dot{p}_{Am}\}$; energy conductance, \dot{v} ; somatic maintenance rate, $[\dot{p}_M]$; allocation fraction to soma, κ ; maturity level at puberty, E_H^p ; and maturity densities E_H^*/L_*^3) relative to other related species in the Add-my-pet collection [20]. Parameters that scale with size ($\{\dot{p}_{Am}\}$ and E_H^p) are divided by the zoom factor to enable comparisons between different-sized species [16]. The parameter values belonging to the same or closely related species are more similar than those of different species. Other sea turtle species (with AmP dataset references in parentheses) are: *Chelonia* mydas [dataset] 2017, *Lepidochelys kempii* [dataset] 2015, *Natator depressus* [dataset] 2017, and *Dermochelys* coriacea [dataset] 2015. The river turtle *Carettichelys insculpta* [dataset] 2017 (marked with a red asterisk) was included because of similar morphology (flippers).

⁵³⁴ in a food-deprived environment.

A higher value of volume-specific somatic maintenance for MED turtles may be linked to 535 the energetic costs of osmotic regulation related to the higher salinity of the Mediterranean. 536 Loggerhead turtles are excellent osmoregulators [101, 102, 103], but coping with the increased 537 salt load by a more intense secretion of the lachrymal salt glands can lead to dehydration 538 [104] and presumably carries an additional energetic cost reflected in the increased cost of 539 maintenance [105, 106]. While lower salinity does not seem to have adverse effects [107, 108], 540 areas of the Mediterranean with higher salinity may act as a "salinity barrier" sensu Carreras 541 et al. [2], thus restricting NA loggerhead turtles to areas with lower salinity [2, 28]. 542

Maturity levels at birth and puberty (E_H^b and E_H^p , respectively) are metabolic switches that mark transitions between embryo, juvenile, and adult stages independent of size or age ⁵⁴⁵ [16]. The difference between maturity at puberty is striking between the two populations of ⁵⁴⁶ loggerhead turtles: MED loggerhead turtles need to invest 40% less energy than NA turtles ⁵⁴⁷ to reach puberty, and they need to allocate around 22 kJ d^{-1} less energy to maintain the ⁵⁴⁸ maximum level of maturity (Fig. 7). Less investment into maturation and the corresponding ⁵⁴⁹ lower maintenance is certainly favorable in low-food environments because—all else being ⁵⁵⁰ equal—it leads to earlier sexual maturation and higher reproductive output at the same ⁵⁵¹ body size.

Faster growth and earlier maturation at a smaller size have three main energetic benefits: (i) less cumulative energy invested into growth, (ii) less cumulative energy spent on metabolic maintenance, and (iii) sexual maturation reachable at lower food levels [14]. This strategy is therefore beneficial in resource-poor environments with short periods of food availability as faster growth uses the (temporarily) available resources to rapidly increase in size, and smaller size at sexual maturity typically requires less energy to complete a reproduction cycle [15, 109].

Interestingly, the ratio of the maturity level and the predicted structural volume, i.e., maturity density [110, 111, this issue] at a specific life stage transition is strikingly similar for both populations (Fig. 8). A correlation between maturity density at birth and maturity density at puberty seems to exist for all species of turtles (last panel in Fig. 8), and all animal taxa in the Add-my-pet collection [111, this issue]. Perhaps species-specific maturity densities rather than maturity levels drive metabolic switches (for more details see section 1.1.4. of the online comments file, [112], pages 5-8).

The apparent success of our modeling approach to put the size dimorphism exhibited by 566 the two studied populations in an evolutionary context points to a deeper connection between 567 DEB theory and evolution. When individuals of a certain species move to expand the range 568 of their species, they are likely to be less-than-optimally adapted to the new territory. In such 569 circumstances, natural selection favors those adaptations that help sustain the presence in the 570 new territory, implying that over an evolutionary time scale, even the parameters of a DEB 571 model become dynamical variables. Future applications of DEB theory—e.g., in relation to 572 conservation efforts [96]—should perhaps consider the kind of a modeling approach that can 573 capture the dynamics of such variables. Jusup et al. [97] have made an early proposal in this 574 direction, but the usefulness of their proposal is yet to be seen. 575

576 5. Conclusion

⁵⁷⁷ We demonstrate that the size dimorphism between the two studied populations of log-⁵⁷⁸ gerhead turtles cannot be explained solely by the difference in environmental conditions, but requires a population-specific description. An implication is—also consistent with the genetic separation between the two populations [2, 3, 4]—that adaptations to the new environment drive the evolution of metabolic traits over sufficiently long time scales.

⁵⁸² We find that three DEB primary parameters capture most of the differences in growth, ⁵⁸³ maturation, and reproduction between North Atlantic (NA) and Mediterranean (MED) log-⁵⁸⁴ gerhead turtles: $[\dot{p}_M]$, $\{\dot{p}_{Am}\}$, and E_H^p . Higher somatic maintenance $([\dot{p}_M])$ and lower cumula-⁵⁸⁵ tive investment to maturity (E_H^p) of the MED population are interpreted here as adaptations ⁵⁸⁶ to environmental factors such as high salinity, which increases the energetic cost of salt gland ⁵⁸⁷ activity, and low food availability, which favors sexual maturation at a smaller size. More ⁵⁸⁸ specifically, we find that:

(i) MED loggerhead turtles grow and mature faster than their NA cousins when in the Mediterranean Sea due to a faster assimilation (i.e., larger $\{p_{Am}\}$), but reach smaller ultimate size due to lower food availability (lower f) and higher somatic maintenance (larger $[\dot{p}_M]$);

(ii) MED loggerhead turtles become sexually mature at a smaller size than their North Atlantic cousins due to a lower cumulative investment to maturation (lower E_H^p). The ratio of the invested energy to the structural volume, dubbed maturity density, might indicate the corresponding maturity level for a metabolic switch when sizes at life events drastically differ between genetically distinct populations of a same species.

(iii) The smaller ultimate size of MED loggerhead turtles is a consequence of low food availability in the Mediterranean Sea, but the reproductive output is comparable to that of NA individuals because more energy is available for reproduction due to a lower maturity maintenance (proportional to E_H^p).

(iv) A combination of lower food assimilation potential and higher costs of maturity maintenance in fully grown NA loggerhead adults offers a metabolic explanation for the observation that NA females do not nest in the Mediterranean.

604 6. Acknowledgements

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Appendix A. Size data on North Atlantic (NA) and Mediterranean (MED) log gerhead turtles, and additional results from statistical analyses

All data used for the comparison is listed in Tables A.4 to A.6. The data was either 614 available as a dataset, or reported as a mean with a range and/or standard deviation (SD). 615 For the MED population, substantially fewer datasets were available. The data for wild 616 populations was mostly reported as a mean with a range and/or standard deviation. In 617 addition, we used a previously unpublished dataset from a rearing and reproduction program 618 in Marineland Antibes (France). The dataset contained measurements of weight at hatching 619 and weekly weight increase up to the age of 2 years for 21 juveniles, and measurements of 620 size and weight of two females reproducing in captivity. This data cannot be analyzed as a 621 part of the wild MED population even though the adults were obtained (and reproduce) in 622 the Mediterranean – the captive reared individuals experience considerably more favorable 623 conditions than the turtles in the wild, and are therefore expected to have a better condition 624 index and produce heavier hatchlings [16]. Consequently, the data from the reproduction 625 program was analyzed separately. 626

For the analysis of the size at hatching, data reported at the moment of hatching (leaving 627 the egg) was pooled together with the data reported for the moment of emergence (leaving 628 the nest). This was justified as the length does not significantly change between hatching 629 and emergence. The minor decrease in wet weight between hatching and emergence (due to 630 dehydration, Bennett et al. [113]) was thus ignored. Length and/or wet weight are mostly 631 reported at emergence for the field incubated nests (e.g. [37]), and at hatching for the 632 laboratory incubated nests (e.g. [38]); pooling the data for analysis increased the sample size 633 and aided the comparison. 634

⁶³⁵ By focusing on "size at nesting" rather than "size at puberty" we were able to use literature ⁶³⁶ data for nesting turtles, which is mostly reported in such a way that data describing first-time ⁶³⁷ nesters cannot be distinguished from that describing experienced nesters.

Table A.4: Egg diameter and wet weight – overview of the data used in the analysis. The range [minimum and maximum] and/or the standard deviation (SD) of the sample is given in the square brackets and parenthesis, respectively, and the number of data points in italicized font (N) where the information was available. Data sources (ds) are indicated next to each data set.

Diameter (cm)				Mass (g)			
North Atlantic	ds	Mediterranean	ds	North Atlantic	ds	Mediterranean	ds
4.25 (0.14), 48	1	$3.76\ (0.142),\ 23$	1	42.58(1.78), 45	2	27.6 (3.1), 23	1
		3.74 (0.196) ^a [3.29-3.96], <i>12</i>	3^1	39.4 (3.8), 48	1	$32.4 (3.52)^{a} [26.4-38.6], 12$	3^1
		$3.8-3.86^{\rm b} (0.133)^{\rm a}$ [3.49-4.02], 13-26	3^1			$26.9-31.9^{\rm b} (3.42)^{\rm a}$ $[22.8-36.5], 13-30$	3^1
		$3.61-3.87^{\rm b} (0.390)^{\rm a}$ [2.7-4.26], 12-30	3^2			$29.8 (3.10)^{a} [23-35.4],$ 45	3^3
		$3.67 (0.223)^{a}$ [3.3-4.19], 45	3^2			$27.5 (3.43)^{a}$ [15.9-36.5], 173	3^2
		$3.45 (0.174)^{a} [3.1-3.5],$ 15	3^3			30.48 (1.62),10	4
		$3.7 (0.200)^{a} [3.3-4.1],$ 65	3^3			30.21 (1.65), 10	4
		$\frac{4.04 - 4.21^{\rm b} \ (0.232)^{\rm a}}{[3.7 - 4.5], \ 5 - 8}$	3^3			30.31 (1.79), 10	4

^a standard deviation calculated following Hozo et al. [51], no distribution assumed

 $^{\rm b}$ data for more than one season reported together; for our analysis we used a non-weighted mean of means and an average sample size

Data source key: {1} Tiwari and Bjorndal [5] (Greece and Florida), {2} Ackerman [50] (Florida),

{3} Margaritoulis et al. [1] and references therein – sample sites: ¹Cyprus; ²Greece; ³Turkey; {4} Reid et al.
 [38] (Greece)

Table A.5: Hatchling length (SCL) and wet weight – overview of the data used in the analysis. The range [minimum and maximum] and/or the standard deviation (SD) of the sample is given in the square brackets and parenthesis, respectively, and the number of data points in italicized font (N) where the information was available. Data sources (ds) are indicated next to each data set. Data from the reproduction program is included as a separate group, with SCL at hatching unknown.

Length (cm)			Mass (g)				
North Atlantic	ds	Mediterranean	ds	North Atlantic	ds	Mediterranean	ds
4.48 (0.159) [4.13-5.0], <i>110</i>	1^1	$\frac{4 \ (0.407)^{a}}{2064} \ [2.49-4.93],$	3^{1}	22.08 (1.49) 41	2	$15.3 (2)^{a} [9.4-21.4],$ 1482	3^{1}
$\begin{array}{c} 4.56 \ (0.141) \\ [4.1-4.99], \ 129 \end{array}$	1^{2}	$\begin{array}{l} 4.03\text{-}4.15^{\mathrm{b}} \ (0.150)^{\mathrm{a}} \\ [3.6\text{-}4.5], \ 180\text{-}325 \end{array}$	3^1	$18.03 (1.33) [14-21], \\ 110$	1^1	$\begin{array}{l} 15.9\text{-}16.7^{\mathrm{b}} \ (1.583)^{\mathrm{a}} \\ [12\text{-}21.5], \ 180\text{-}325 \end{array}$	3^1
$\begin{array}{c} 4.43 \ (0.134) \\ [4.09-4.71], \ 114 \end{array}$	1^1	4.04 (0.7), 20	3^2	19.87 (1.43), <i>129</i>	1^2	16.74 (0.82), 10	4
4.45 (0.17) [3.96-4.76], <i>108</i>	1^1	3.98 (0.407) ^a [2.8-4.5], <i>302</i>	3^3	18.71 (2.18) [14-24.5], 114	1^1	16.72 (1.02), 10	4
$\begin{array}{c} 4.42 \ (0.156) \\ [3.87-4.7], \ 106 \\ 4.6 \ (0.11) \ [4.2,4.0] \end{array}$	1^1	$\begin{array}{l} 3.91 \ (0.150)^{\rm a} \\ [3.6-4.2], \ 37 \end{array}$	3^3	18.75 (1.743) [14-22], 108 10.60 (1.743)	1^1	16.59 (0.9), 10	4
4.6 (0.11) [4.3-4.9], 120	5	4.29 (0.09), 10	4	$18.68 (1.743) \\ [14-22.5], 106 \\ 10.0 (1.22) \\ \$	1^1		
		4.24 (0.1), 10	4	19.8 (1.33) [15.3-22.4], <i>120</i>	5		
		4.22 (0.1), 10	4				

^a standard deviation calculated following Hozo et al. [51], no distribution assumed

^b data for more than one season reported together; for our analysis we used a non-weighted mean of means and an average sample size

Data source key: {1} Data partially published in Stokes et al. [37] – sample sites: ¹Florida, ²South Carolina; {2} Ackerman [50] (sample site: Florida); {3} Margaritoulis et al. [1] and references therein – sample sites: ¹Cyprus, ²Greece, ³Turkey; {4} Reid et al. [38] (Greece); {5} Reich et al. [114] (Florida); Additional data was available from the Marineland (Antibes, France) rearing facility : 21.02 (SD 5.8), range [16.7-37.5], N=21 – data could not be approximated by a normal distribution and was analyzed separately.

Table A.6: Adult length (SCL) and wet weight – overview of the data used in the analysis. The range [minimum and maximum] and/or the standard deviation (SD) of the sample is given in the square brackets and parenthesis, respectively, and the number of data points in italicized font (N) where the information was available. Data sources (ds) are indicated next to each data set.

Length (cm)				Mas	s (g)		
North Atlantic	ds	Mediterranean	ds	North Atlantic	ds	Mediterranean	ds
90.9 (4.9) [82-103], <i>84</i>	1	79.43 (4.4) [74.31-84.37], <i>14</i>	2	118.2 (17.5) [89.7-170.9], <i>47</i>	1	$67.26 (8.625)^{a}$ [52.5-87], 40	3
92.3 (5.6) [81-110], 110	1	76.8-80.1 ^b $(6.799)^{a}$ [63.5-87], 11-15	4^1	116.3 (17.1) [71.7-148.9], 93	1		
92.01 (5.34) [78.89-104.47], <i>102</i>	1*	78.6-79.1 ^b (7.250) ^a [66-95], <i>13-97</i>	4^{1}	114.7 (20.3) [79.6-180.7], <i>121</i>	1		
92.5 (2), [85-98], 13	5^1	$78.3-79.2^{\rm b} (3.583)^{\rm a} [66-95], 195-343$	4^1	$\begin{array}{l} 93.27 \ (22.64) \\ [46.67\text{-}153.74], \ 56 \end{array}$	6*		
92.4 (0.8) [80.5-107], 52	5^{2}	78.7 $(6.273)^{a}$, [62.3-83.2], 9	4^{2}				
93.1 (1) [83-105], 137^3	5	$73.1 (5.925)^{a} [60.2-83.9], 49$	4^3				
90.9 (5) [76.80-100.28], <i>51</i>	2	$73.2 \ (5.375)^{\rm a} \\ [66-87.5], \ 22$	4^{3}				
94.3 (5.5) [83.8-106.7], <i>41</i>	7	72 $(7.250)^{a}$ [58-87], 58	4^{3}				
95.1 (4.8) [80.7-107.4], <i>84</i>	7						
94.73 (5.29) [80.72-107.34], <i>112</i>	7*						
89.59 (6.56) [80.56-105.04], <i>56</i>	6*						

^a standard deviation calculated following Hozo et al. [51], normal distribution assumed

^b data for more than one season reported together; for our analysis we used a non-weighted mean of means and an average sample size

Data source key: [1] Ehrhart and Yoder [69], *digitalized from Figure 3 (sample site: Florida); [2] Tiwari and Bjorndal [5] (Greece and Florida); [3] Groombridge [11] (Greece); [4] Margaritoulis et al. [1] and references therein – sample sites:¹Greece, ²Lybia, ³Turkey; [5] Stoneburner [31] (sample sites: ¹North Carolina, ²Georgia, ³Florida); [6] Wabnitz and Pauly [68], *digitalized from Figure 1, only used samples from North Atlantic for loggerhead turtles longer than 80.5 cm SCL (sample sites: Virginia, Florida); [7] Byrd et al. [70], *digitalized from Figure 3 (sample sites: South Carolina); Additional data was available from the Marineland (Antibes, France) rearing facility – individual values for two females: SCL = 70 cm and 80 cm, W = 86.6 kg and 100.3 kg – data was analyzed separately.

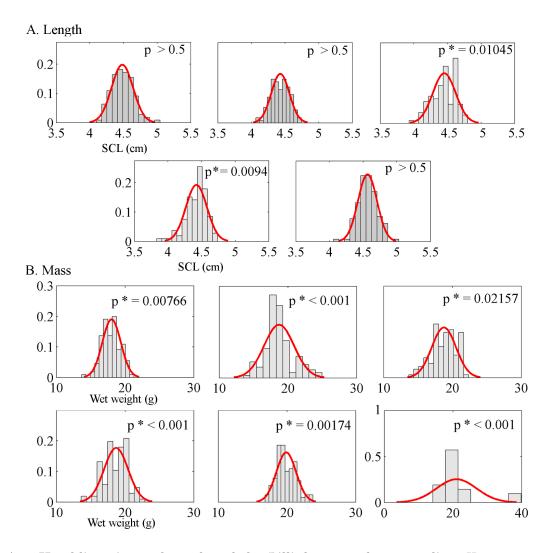


Figure A.9: Hatchling size and results of the Lilliefors test for normality. Histograms represent data and the solid line is the model of normal distribution based on the mean and standard deviation of the corresponding sample plotted across a range of three standard deviations from the mean. Even though some of the samples significantly differ from the normal distribution (indicated with a star next to the p-value), histograms of some datasets visually resemble a normal distribution. Panel a. Length. Panel b. Mass. All data for length and most data for mass are courtesy of L. Stokes (partially published in Stokes et al. [37]) and are from the NA population; data for mass include an additional previously unpublished dataset from Marineland (Antibes, France) rearing facility.

Lilliefors test. Test rejected the null hypothesis of that sizes at hatching are sampled from a normal distribution (Lilliefors test, p < 0.05) for two out of five length datasets and all six tested mass datasets. However, histograms suggest that the discrepancies are driven by just a few outliers that skew the distribution, which would otherwise resemble the normal one (Figure A.9, panels a and b). We therefore chose to assume the normal distribution for all datasets except those describing the captive reared hatchings, for which the discrepancies from the normal distribution were substantial. The size at nesting seems to follow the normal distribution even more closely: the test could not reject the null hypothesis (Lilliefors test, p > 0.05) for none of the tested length datasets (see also Figure A.10).

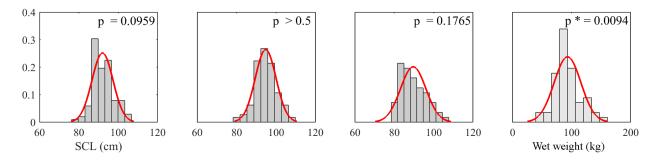


Figure A.10: Adult size and results of the Lilliefors test. Histograms represent data and the solid line is the model of normal distribution based on the mean and standard deviation of the corresponding sample plotted across a range of three standard deviations from the mean. The mass sample (last panel) significantly differs from the normal distribution (indicated with a star next to the p-value). Data sources: [68, 69, 70].

Welch's F-test and unequal variances t-test. The hypothesis that all samples de-647 scribing the same life stage came from the same statistical population was rejected (Welch's 648 F-test p > 0.05, Figure 1), supporting the hypothesis that samples from the two popula-649 tions were different. Statistically even samples collected on neighboring beaches were not 650 from the same population [37]. However, we grouped samples into NA and MED groups 651 for the purpose of further analysis because: (i) visual analysis of probability density plots 652 suggested a grouping of such type (Fig. 1); (ii) pairwise Welch's t-tests suggested that the 653 size data assigned to the NA population were significantly larger than those assigned to the 654 MED population in almost all cases (Fig. A.11); and (iii) grouping enabled inter-population 655 comparison on a more general level. 656

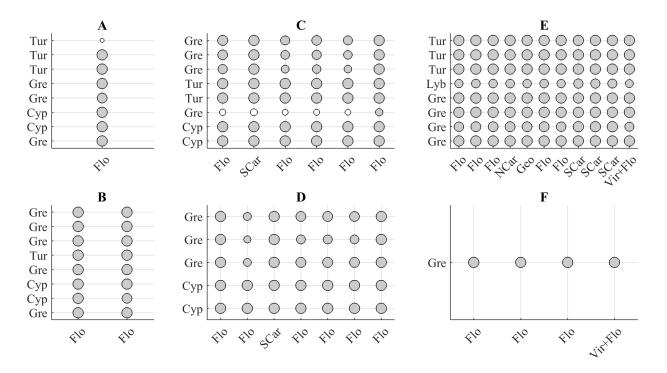


Figure A.11: Results of the pairwise Welch's t-test for sizes at the analyzed life events. Full circles denote the samples are significantly different (p < 0.05) with the size of the circle proportional to statistical significance. **Egg** diameter (panel A) and mass (panel B); **Hatchling's** length (panel C) and mass (panel D) at hatching, and **adult female's** length (panel E) and mass (panel F) at nesting. Sampling location are indicated at the vertical axis for the MED population (Tur – Turkey, Gre – Greece, Cyp – Cyprus, Lyb – Lybia), and horizontal axis for the NA population (Flo – Florida, SCar – South Carolina, NCar – North Carolina, Vir – Virginia).

⁶⁵⁷ Appendix B. Overview of the standard Dynamic Energy Budget (DEB) model ⁶⁵⁸ for loggerhead turtles

The standard DEB model follows changes over time of three main state variables (marked S1-S3 in Fig. B.12), and an auxiliary variable (R1 in Fig. B.12, present only in adults). Temperature is included as an effect on all rates [16]. Food availability is included as the scaled functional food response, i.e., a saturating function denoting the feeding rate as a fraction of the maximum for an individual of the same size [16, 95].

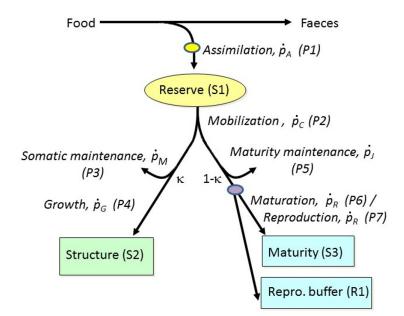


Figure B.12: Schematic representation of the standard DEB model. Dynamics of the state variables are determined by energy flows universally denoted \dot{p}_* (unit Jd^{-1}). For dynamics of state variables see Table B.7, for process descriptions and equations Table B.8, and for parameters of the model Table 2. Food availability is quantified by the scaled functional response, f (i.e., f = 1 under unlimited food availability and f = 0 when food is unavailable, see p. 32 of [16] for details.) Circles mark metabolic switches: birth (yellow circle) is the onset of feeding, whereas puberty (purple circle) is the onset of investement into reproduction. Mobilized reserve is partitioned according to the κ -rule: fixed fraction κ is allocated to satisfy the organism's somatic needs (somatic maintenance and growth), whereas the rest is allocated to maturity maintenance and maturation (before puberty) or reproduction (after puberty).

Table B.7: State variables of the standard DEB model. S1-S3 in the "Notes" column refer to the
corresponding compartment in Fig. B.12. For energy flows see Table B.8, for parameter descriptions and
values Table 2.

State variable	Equation	Note
Reserve	$\frac{dE}{dt} = \dot{p}_A - \dot{p}_C$	S1
Structure	$\frac{dE}{dt} = \dot{p}_A - \dot{p}_C$ $\frac{dL}{dt} = \frac{1}{3L^2} \frac{\dot{p}_G}{[E_G]}$	S2
Maturity	$\frac{dE_H}{dt} = \dot{p}_R,$ if $E_H < E_H^p$ $\frac{dE_H}{dt} = 0,$ otherwise	S3; E_H^p is maturity at puberty marking the beginning of the adult stage (see Fig. B.12. In this stage, investment into maturation gets redirected into reproduction: $\frac{dE_R}{dt} = \dot{p}_R$ (auxiliary variable R1 in Figure B.12)

Table B.8: **Energy flows** appearing in the system of equations for state variables (Table B.7), modified from Marn et al. [19]. All model parameters are conveniently summarized in Table 2.

Process	Equation	Note
Assimilation	$\dot{p}_A = \{\dot{p}_{Am}\}fL^2$	P1; The fraction of the daily feed ration that gets fixed into reserve, where $\{\dot{p}_{Am}\}$ (unit J cm ⁻² d ⁻¹) is the surface area-specific maximum assimilation rate and f is the scaled functional response equivalent to the ratio of the actual and the maximum feeding rate of an individual.
Mobilization	$\dot{p}_C = E(\dot{v}/L - \dot{r})$	P2; The flow of energy mobilized from reserve to power metabolic processes, where parameter \dot{v} (unit d ⁻¹) is the energy conductance and, for $[E] = E/L^3$, the specific growth rate is $\dot{r} = \frac{[E]\dot{v}/L - [\dot{p}_M]/\kappa}{[E] + [E_G]/\kappa}$. Here, $[\dot{p}_M]$ (unit J cm ⁻³ d ⁻¹) is the volume-specific somatic maintenance rate.
Somatic maintenance	$\dot{p}_M = [\dot{p}_M]L^3$	P3; The flow of mobilized reserve energy needed to maintain the structure of given size L^3 .
Growth	$\dot{p}_G = \kappa \dot{p}_C - \dot{p}_M$	P4; The flow of mobilized reserve energy invested into the increase of structure after satisfying the somatic maintenance needs.
Maturity maintenance	$\dot{p}_J = \dot{k}_J E_H, E_H \le E_H^p$	P5; A flow (analogous to somatic maintenance) that quantifies the mobilized reserve energy necessary to maintain the current level of maturity. Parameter \dot{k}_J (unit d^{-1}) is called the maturity maintenance rate coefficient.
Maturation	$\dot{p}_R = (1 - \kappa)\dot{p}_C - \dot{p}_J$	P6; The flow of mobilized reserve energy towards increasing the level of maturity (E_H) , after satisfying the maturity maintenance, \dot{p}_J .
Reproduction	\dot{p}_R	P7; At the onset of the adult stage when the level of maturity reaches E_H^p , the organism starts to invest energy into reproduction instead of maturation. Hence, reproduction starts and maturity stops increasing.

We first used the parameter set for the North Atlantic (NA) population of loggerhead 664 turtles $(pars_{NA})$ to run the model and get predictions for environmental characteristics de-665 scribing the Mediterranean Sea (Fig B.13, see Section 2.2 for more details). We compared 666 predictions (life history traits, growth curves, reproduction output etc.) to observations for 667 Mediterranean (MED) loggerhead turtles to calculate the goodness of fit [21, 58, this issue]. 668 In the second step, we used the observations (on life history traits, growth, and reproduc-669 tion) in addition to $'pars_{NA}$ ' and environmental characteristics, and with the covariation 670 routines obtained a new parameter set $('pars_{MED}')$ and model predictions. The covariation 671 routines (integrated into DEBtool [61]), use the Nelder-Mead method and symmetric-bound 672 loss function for parameter estimation [57, 58, this issue]. 673

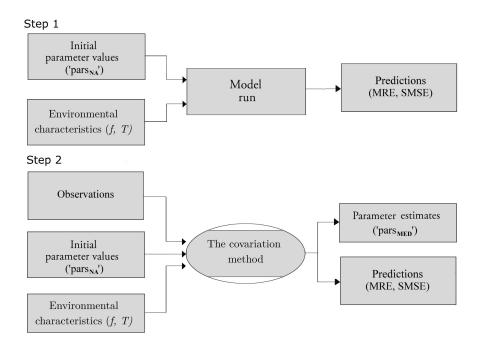


Figure B.13: Schematic representation of the two steps used in the mechanistic approach. Parameter set ' $pars_{NA}$ ' was estimated with the covariation method previously using only data for North Atlantic (NA) loggerhead turtles (Marn et al. [19] and AmP *Caretta caretta* [dataset] [71]). Parameter set ' $pars_{MED}$ ' pertains to the Mediterranean (MED) loggerhead turtles. Goodness of fit to data is expressed as mean relative error (MRE) and symmetrical mean squared error (SMSE). See Section 2.2 for more details

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