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Regional environmental drivers of Kemp’s ridley sea turtle somatic growth variation

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23 **Abstract**

24 Many environmental processes influence animal somatic growth rates. However,
25 elucidating specific drivers of somatic growth variation has been challenging for marine
26 megafauna. Using a 20+ year dataset of somatic growth generated through skeletochronology,
27 we evaluated the relationship between multiple region-wide environmental factors—the
28 *Deepwater Horizon (DWH)* oil spill, increasing population density, climate variability—and age-
29 and region-specific Kemp’s ridley sea turtle (*Lepidochelys kempii*) somatic growth. We observed
30 significant, multi-year reductions in mean oceanic (age 0) and small neritic (age 2–5) juvenile
31 growth rates beginning in 2012 for turtles stranded along the U.S. Gulf of Mexico (GoM) and
32 Atlantic Coasts, which resulted in a reduction in mean size-at-age. We hypothesize this growth
33 decline is related to long-term deleterious effects of the *DWH* oil spill on benthic and oceanic
34 food webs in the GoM. Additionally, regional climate indices were strongly correlated with
35 oceanic juvenile growth with a 2-yr lag (cross-correlation = -0.57 to 0.60), whereas GoM small
36 neritic juvenile growth was strongly related to population abundance metrics. Generalized
37 additive models that included all examined environmental covariates indicated that the drivers of
38 the 2012 growth rate decline had the strongest effect on Kemp’s ridley growth rates between
39 1995 and 2015, but that additive or synergistic effects of both climate variability and changing
40 population abundance are likely for certain life stages. Continued collection of sea turtle humeri
41 is needed to further clarify mechanisms underpinning the observed growth patterns given the
42 coincidental timing of changes in environmental parameters examined herein.

43

44 **Keywords:** somatic growth rates, *Lepidochelys kempii*, density-dependence, climate effects,
45 skeletochronology, environmental covariates, Gulf of Mexico

46 **Introduction**

47 A suite of natural and anthropogenic stressors have reshaped marine ecosystems over the
48 past century through cascading effects on animal populations and the habitats they occupy
49 (Halpern et al. 2008; Rocha et al. 2014; McCauley et al. 2015). Numerous studies have
50 characterized single stressor effects on marine species, but fewer have examined species
51 response to cumulative or integrative effects of multiple environmental stressors, particularly in
52 long-lived, higher order marine megafauna (Crain et al. 2008; Bjorndal et al. 2013). As the
53 population dynamics of long-lived species are highly sensitive to small changes in demographic
54 rates (Heppell et al. 2000), increasing insight into environmental effects on growth, survival, and
55 reproduction may help improve understanding of population and community dynamics, and
56 ultimately aid the development of conservation and management strategies for protected species.
57 Moreover, disentangling the relative influence of myriad environmental stressors on animal
58 populations and ecosystems is essential to predicting future ecosystem response to perturbation.
59 Sea turtles provide an ideal system to investigate the influence of multiple environmental
60 phenomena on demographic rates because most species retain annual records of somatic growth
61 in their humerus bones, similar to growth rings in trees and otoliths in fish, that can be collected
62 from dead stranded turtles (Avens & Snover 2013), and, as ectotherms, their growth rates are
63 highly influenced by environmental conditions.

64 The critically endangered Kemp's ridley sea turtle (*Lepidochelys kempii*) is a particularly
65 appealing model species to evaluate environmental drivers of somatic growth rates. First,
66 humerus bones have been collected from dead stranded turtles since the early 1990s (Snover and
67 Hohn 2004; Avens et al. 2017), providing a unique sample set for growth analysis. Second, their
68 global distribution is largely restricted to the Gulf of Mexico (GoM) and U.S. Atlantic (Musick

69 and Limpus 1997), areas that are experiencing rapid environmental change including a climate-
70 driven ecological regime shift in the 1990s (Sanchez-Rubio et al. 2011; Karnauskas et al. 2015)
71 and the 2010 *Deepwater Horizon (DWH)* oil spill (DWH NRDA Trustees 2016). The current
72 understanding of Kemp's ridley life history suggests juveniles that reside in benthic GoM and
73 U.S. Atlantic Coast habitats are geographically isolated from one another, with some proportion
74 of turtles entering the U.S. Atlantic Coast at age 1 or 2 following the GoM oceanic life stage and
75 not returning to the GoM until maturity (Putman et al. 2013, Caillouet et al. 2016, Avens et al. *in*
76 *review*). This geographic isolation of two components of the population provides a natural
77 experiment to examine *DWH* oil spill effects on sea turtle growth rates and potentially separate
78 them from other region-wide environmental stressors. Lastly, the Kemp's ridley population grew
79 exponentially (12–16% per year) through the 1990s and 2000s following decades of successful
80 conservation and management (NMFS and USFWS 2015). This, combined with a robust record
81 of nest and hatchling production for nearly the entire species, provides the opportunity to
82 evaluate density dependent effects on their somatic growth rates (Caillouet et al. 2016, 2018).

83 Environmental impacts of the *DWH* oil spill and impact mitigation efforts were
84 unprecedented in their spatiotemporal and ecological scale (DWH NRDA Trustees 2016; Beyer
85 et al. 2016; Berenshtein et al. 2020). Negative effects of the *DWH* oil spill on somatic growth
86 rates have been documented in a wide range of fish and invertebrate species (e.g., Rozas et al.
87 2014; Brown-Peterson et al. 2016; Herdter et al. 2017; Perez et al. 2017), but impacts on marine
88 megafauna demographic rates are less understood. However, long-term impacts remain a
89 significant concern given the continued deterioration of the health of GoM bottlenose dolphins
90 and the clear decadal impacts of the 1989 *Exxon Valdez* oil spill on marine ecosystems and
91 animal demographic rates (Peterson et al. 2003; Kellar et al. 2017). Immediate effects on sea

92 turtle survival and physiology are well-documented, but otherwise much remains unknown about
93 their response to this anthropogenic disturbance (McDonald et al. 2017; Mitchelmore et al. 2017;
94 Stacy et al. 2017; Wallace et al. 2017). Sublethal or indirect effects of the *DWH* oil spill on sea
95 turtle health may be responsible for a general decline in nutritional condition of stranded sea
96 turtles since 2012 and a reduction in juvenile Kemp's ridley growth rates in Mississippi since
97 2010 (Stacy 2015; Coleman et al. 2016).

98 Following decades of conservation and management, the abundance of all Kemp's ridley
99 life stages grew rapidly between 1990 and 2009 (Heppell et al. 2004; NMFS and USFWS 2015).
100 Unexpectedly, annual nest counts have fluctuated widely since 2010 and one hypothesis is that
101 density dependent processes may be acting on the population (Gallaway et al. 2016; Caillouet et
102 al. 2016, 2018). While the current population is less than 10% of its estimated historic size
103 (Bevan et al. 2016), long-term alteration and degradation of GoM ecosystems, including
104 reductions in important food resources (e.g., blue crab *Callinectes sapidus*; VanderKooy 2013),
105 may have lowered the potential carrying capacity of the GoM for sea turtles and other marine top
106 predators (Heppell et al. 2007; Caillouet 2014). Most support for this hypothesis is derived from
107 analyses of the species' nesting trends (Gallaway et al. 2016; Caillouet et al. 2016, 2018;
108 Kocmoud et al. 2019), which are confounded after 2010 with unknown effects of the *DWH* oil
109 spill, and the observation of increasing breeding intervals for Kemp's ridleys nesting in Texas
110 from 2008 to 2016 (Shaver et al. 2016). However, other environmental factors, such as colder
111 temperatures on the foraging grounds during the winter of 2009–2010 (Lamont and Fujisaki
112 2014; Gallaway et al. 2016), may underpin this change in breeding interval and additional
113 investigations are needed to evaluate whether density dependent processes are influencing
114 Kemp's ridley demographic rates.

115 Climate variability is a primary driver of spatiotemporal variability in ocean productivity,
116 and abrupt changes in climate forcing often precipitate ecological regime shifts (Rocha et al.
117 2014). Within the North Atlantic Ocean, an ecological regime shift occurred in the late-1990s as
118 a result of an abrupt warming of the ocean that coincided with one of the strongest El Niño
119 events on record as well as a shift from the cool to warm phase of the Atlantic Multidecadal
120 Oscillation (Sanchez-Rubio et al. 2011; Luczak C. et al. 2011; Reid and Beaugrand 2012;
121 Beaugrand et al. 2013; Karnauskas et al. 2015). This late-1990s regime shift has been linked to
122 reduced blue crab productivity in the GoM (Sanchez-Rubio et al. 2011), an important food
123 source for sea turtles, as well as declining growth rates in loggerhead (*Caretta caretta*), green
124 (*Chelonia mydas*), and hawksbill (*Eretmochelys imbricata*) sea turtles (Bjorndal et al. 2013,
125 2016, 2017). Similar declines in growth were observed in large juvenile and adult Kemp’s
126 ridleys in the GoM from 1988 to 2009 and small juveniles from 2004 to 2009 (Avens et al.
127 2017), although links to climate variability have yet to be evaluated.

128 Here we examined temporal trends in juvenile Kemp’s ridley sea turtle somatic growth
129 rates using a 20+ year dataset generated through skeletochronology. The primary objective of
130 this study was to quantify the relative influence of multiple regional environmental stressors—
131 the *DWH* oil spill, increasing population density, climate variability—on sea turtle growth rates.
132 We developed and tested a suite of hypotheses related to the differential effect of these factors
133 that are outlined here and in Figure 1. Given significant degradation of offshore and nearshore
134 habitats in the GoM following the 2010 *DWH* oil spill and the observed decline in GoM-stranded
135 turtle nutritional condition after 2012 (Stacy 2015; Beyer et al. 2016), we predicted that Kemp’s
136 ridley growth rates would decline following the *DWH* oil spill for both oceanic and neritic
137 juveniles. We specifically predicted that this change would occur beginning in 2010 because

138 annual Kemp's ridley skeletal growth begins in spring, coincident with the timing of the *DWH* oil
139 spill. Importantly, we predicted Atlantic turtle growth rates would not change after 2010 given
140 their spatial isolation from the *DWH* oil spill. We predicted that density-dependent effects, if
141 present, would result in declining growth rates beginning in the mid- to late-2000s, when
142 population growth was the highest (NMFS and USFWS 2015). We expected density dependent
143 effects would primarily manifest in small juvenile life stages in the GoM as they have the fastest
144 growth rates and experience the greatest competition with conspecifics for resources due to their
145 size and relative inexperience. We assumed Atlantic Kemp's ridley are not strongly influenced
146 by intraspecific population density due to their relatively low abundance. Lastly, we predicted
147 that climate effects would cause declining growth rates across all Kemp's ridley life stages and
148 habitats beginning in the late-1990s in response to a regional regime shift as observed in other
149 western North Atlantic sea turtle species (Bjorndal et al. 2016, 2017).

150

151 **Materials and Methods**

152 *Sample collection and processing*

153 Front flippers were collected from Kemp's ridleys that stranded on U.S. beaches by
154 participants of the Sea Turtle Stranding and Salvage Network (Texas to Massachusetts, 1991 to
155 2017). Samples were obtained from turtles that either stranded dead or stranded alive but were
156 later euthanized. Stranding location, date, and carapace length were recorded at the time of
157 stranding (see Tables 1 and S1 for summary). Carapace length was measured as straightline
158 (SCL) or curved (CCL) carapace length, notch to tip. In cases where only CCL was recorded,
159 CCL was converted to SCL as described by Avens et al. (2017). This study utilizes and extends
160 the growth datasets presented in Avens et al. (2017) ($n = 333$ turtles, GoM) and Snover et al.

161 (2007) ($n = 144$, Atlantic) to include growth histories obtained from a total of 784 turtles
162 stranded along the U.S. GoM Coast and 451 turtles stranded along the U.S. Atlantic Coast. We
163 assume that data derived from these strandings are generally reflective of turtles within each
164 region but acknowledge that strandings represent a non-random sampling of the population. The
165 likelihood of a dead turtle stranding is determined by its decomposition rate and drift time and
166 distance which are in turn influenced by ocean current and temperature, factors that vary across
167 space and time. And, the probability of a stranded turtle being observed is influenced by
168 coastline accessibility. Combined, we can assume strandings are biased towards turtles that die
169 nearshore, perhaps skewing the dataset towards younger/smaller life stages that inhabit more
170 shallow marine habitats, and exclude most oceanic stage turtles, though data for this life stage
171 can be retained in the bones of small neritic juveniles.

172 Humerus bones were prepared and histologically processed as described by Avens and
173 Snover (2013) and Avens et al. (2017). Tissue was removed from the humerus bones, which
174 were then boiled and air dried for at least two weeks. A low speed isomet saw (Buehler) was
175 used to cut a 2 to 3 mm thick cross-section from each bone just distal to the deltopectoral muscle
176 insertion scar. Bone sections were fixed and decalcified using Cal Ex II (Fisher Scientific) or
177 10% neutral buffered formalin followed by RDO (Apex Engineering Corporation) and thin
178 sectioned to 25 microns using a freezing-stage microtome (Leica) or cryostat (Thermo Scientific
179 Microm HM 550). Thin sections were stained using diluted Ehrlich's hematoxylin, mounted onto
180 microscope slides in 100% glycerin, and imaged using a digital camera fitted to a compound
181 microscope. Growth mark analyses were performed using image analysis software (Olympus
182 Microsuite and cellSens) and Adobe Photoshop (Adobe Systems). Two or three readers (of L.
183 Avens, L. R. Goshe, M. Ramirez, M. Snover) independently analyzed the digital bone images to

184 determine the number and placement of lines of arrested growth (LAGs), which delimit the outer
185 edges of each skeletal growth mark (Snover and Hohn 2004), followed by a joint assessment to
186 reach consensus. Once consensus was reached, total humerus section diameter and the diameter
187 of each LAG were measured.

188

189 *Age and growth rate estimation*

190 Previous analyses validated annual LAG deposition in Kemp's ridley humerus bones
191 (Snover and Hohn 2004; Avens et al. 2017), allowing for characterization of age at stranding
192 through skeletochronology. Kemp's ridleys deposit a unique first-year growth mark, or
193 "annulus," that differs from subsequent marks (Snover and Hohn 2004). For bones where the
194 annulus was visible, an initial age estimate was determined directly from LAG counts. However,
195 bone resorption results in the loss of internal LAGs as sea turtles age (Zug et al. 1986),
196 preventing the direct assessment of turtle age in larger individuals where the annulus has been
197 resorbed. Therefore, for turtles where the annulus was not visible, a correction factor was
198 developed based on the relationship between LAG numbers and diameters from known age
199 individuals to estimate the number of LAGs lost to resorption for each bone (Parham and Zug
200 1997). An initial age estimate was then generated by adding the estimated number of resorbed
201 LAGs to the number of visible LAGs. A final age estimate at stranding was made by adjusting
202 initial age estimates to the nearest 0.25 years based on the mean hatch date for the population
203 (June) and individual stranding date. Given that LAG deposition occurs in late winter/early
204 spring and peak hatching for this species occurs during the summer (Snover & Hohn 2004), the
205 first-year growth mark denotes an age of ~0.75 years, the next LAG an age of 1.75 years, and so
206 on. Final age estimates were used to back-assign age estimates to individual LAGs. Similarly, a

207 calendar year was back assigned to each LAG based on the date of stranding.

208 There is a strong allometric relationship between humerus section diameter (HSD) and
209 SCL for Kemp’s ridleys that allows for the back-calculation of body size estimates for
210 measurable LAGs (Snover and Hohn 2004; Avens et al. 2017). We characterized the HSD:SCL
211 relationship for newly processed turtle bones and combined that with the body proportional
212 hypothesis back-calculation technique (BPH; Francis 1990) to estimate SCL for every
213 measurable LAG, adjusted for turtle-specific SCL and HSD at death. Annual somatic growth
214 rates were then calculated by taking the difference between SCL estimates of successive LAGs.
215 In this way, multiple growth rate estimates were generated from each humerus bone (median = 3
216 per turtle, range = 1–8). Growth rate estimates were assigned to the calendar year associated with
217 the LAG that begins each growth interval.

218

219 *Environmental covariates*

220 To investigate environmental drivers of sea turtle somatic growth variation, we evaluated
221 the relative influence of the *DWH* oil spill, changing population density, and climate variability
222 on Kemp’s ridley somatic growth rates. While these stressors are not encompassing of all major
223 environmental phenomena that may affect sea turtle growth rates, they were chosen for this
224 analysis because their potential influence matches the geographic scale encompassed by the
225 somatic growth rate dataset.

226 The relationship between growth and population density was investigated using two
227 population abundance metrics: (1) annual age class-specific abundance estimates obtained from
228 the most recent Kemp’s ridley population model used for status assessment (i.e., model-
229 dependent metric; NMFS and USFWS 2015), and (2) cumulative annual hatchling production

230 from the species' index nesting beach in Tamaulipas, Mexico, which comprises over 85% of
231 nesting activity by the species (i.e., model-independent metric; data sourced from NMFS &
232 USFWS 2015). This species is unique among sea turtles in that nearly its entire annual
233 reproductive output is concentrated on only a handful of beaches in Mexico and South Texas that
234 have been monitored and protected continuously since 1978. This has allowed for the near-
235 complete census of nests laid and hatchlings produced from these beaches annually (NMFS and
236 USFWS 2015). The population model used to derive age-specific abundance estimates is a
237 deterministic age-based simulation model that uses known hatchling production since 1966 to
238 predict the number of nests laid annually (NMFS and USFWS 2015). Model-derived abundance
239 estimates by age-class are only used through 2009 given uncertainties in the cause of post-2009
240 nest count fluctuations—mortality likely increased due to the *DWH* oil spill (Wallace et al. 2017),
241 but other causes have also been proposed (Caillouet 2014; Caillouet et al. 2018; Kocmoud et al.
242 2019), creating substantial uncertainty in the underlying demographic processes for this species
243 after 2009. Trends in population abundance metrics are summarized in Figure S1.

244 To elucidate potential relationships between changes in broad scale climate patterns and
245 Kemp's ridley somatic growth variation, we considered three well-known modes of variability
246 [North Atlantic Oscillation (NAO), Atlantic Multidecadal Oscillation (AMO), and the El Niño
247 Southern Oscillation (ENSO)] that exert strong biophysical control on western North Atlantic
248 Ocean ecosystems (Giannini et al. 2001; Greene et al. 2013; Karnauskas et al. 2015).
249 Collectively, they influence ocean temperature, salinity, mixing, and circulation patterns that
250 affect the productivity, distribution, growth, and survival of animals across all trophic levels
251 (Drinkwater et al. 2003; Edwards et al. 2013; Karnauskas et al. 2015). For the NAO, we used the
252 winter (December to March) NAO index (wNAO) given that the NAO is thought to exert the

253 greatest influence on ocean ecosystems in the boreal winter (Drinkwater et al. 2003). For the
254 ENSO, we used the Multivariate El Niño Southern Oscillation Index (MEI) Version 2, which
255 integrates five meteorological variables: SST, surface air temperature, sea-level pressure, surface
256 zonal winds, surface meridional winds, and Outgoing Longwave Radiation. Monthly AMO and
257 bimonthly MEI data were obtained from NOAA’s Earth System Research Laboratory
258 (<http://www.esrl.noaa.gov/psd/data/climateindices/>) whereas wNAO data were obtained from the
259 National Center for Atmospheric Research (<https://climatedataguide.ucar.edu/climate-data/>).
260 Following Bjorndal et al. (2016, 2017), monthly AMO and bimonthly MEI data were averaged
261 within a calendar year to create an annualized index used in all analyses.

262

263 *Data Analysis*

264 We employed a suite of statistical tools to evaluate the independent and synergistic
265 effects of the *Deepwater Horizon* oil spill, population density, and climate variability on Kemp’s
266 ridley growth rates. Given the retrospective nature of this study, the statistical approach taken
267 was necessarily correlative and we therefore do not conclusively attribute causation. In most
268 cases, analyses were restricted to juvenile growth data—binned by age class (age 0, 1, 2–5, 6–9)
269 to increase statistical power—given that adult turtle growth rate data are poorly represented in
270 the dataset. These age classes align with known ontogenetic differences in somatic growth rates
271 and are similar to those used in age-structured population models (Snover et al. 2007; NMFS and
272 USFWS 2015). Age 0 (ages 0 to 0.75) and 1 (ages 0.75 to 1.75) align with the oceanic life stage
273 but are separated here because growth rates differ between these ages and a fraction of Kemp’s
274 ridleys begin to recruit to neritic habitats at age 1 (Avens et al. *in review*). All other age classes
275 represent neritic life stages, i.e., small neritic juveniles (ages 2–5), large neritic juveniles (age 6–

276 9). As somatic growth rates differ between Kemp’s ridleys that inhabit the U.S. Gulf of Mexico
277 and Atlantic Coast (Avens et al. 2017; Avens et al. *in review*; this study), growth data were
278 analyzed separately for turtles that stranded on beaches in these regions for all age classes but
279 age 0—all age 0 turtles are assumed to occupy the same oceanic habitats in the central GoM .

280 To investigate *DWH* oil spill effects on somatic growth rates we used two primary
281 approaches: growth curve fits and temporal analysis. First, to examine population-level growth
282 response, a family of von Bertalanffy growth functions (VBGFs) were fit to stranding size-at-age
283 data for all turtles stranded before (1993–2009) and after (2011–2016) the *DWH* oil spill using
284 non-linear least-squares regression. Eight models were considered to compare von Bertalanffy
285 growth parameters (L_{∞} , asymptotic average length; K , Brody growth rate coefficient ; t_0 , age
286 when the average length is zero) between both time periods that ranged from including identical
287 parameter estimates for each time period (1 L_{∞} , 1 K , 1 t_0) to including fully unique parameter
288 estimates for each time period (2 L_{∞} , 2 K , 2 t_0), and all model subsets in between (Table 2).
289 Akaike information criterion (AIC) and Akaike weights (w_i) were used to evaluate and compare
290 models (Burnham and Anderson 2002). In addition, given the non-independence of the full
291 growth dataset, VBGFs were fit to measured SCL and estimated age at stranding only,
292 eliminating SCL and age data estimated from growth marks. VBGFs were fit using data from
293 GoM-stranded turtles only; large juvenile and adult Kemp’s ridleys are rare along the U.S.
294 Atlantic Coast and are thus underrepresented in our dataset, preventing the generation of robust
295 Atlantic Kemp’s ridley VBGFs. Growth functions were implemented using the *FSA* (Ogle et al.
296 2018) and *nlstools* (Baty et al. 2015) packages in R (version 3.5.3; R Core Team 2019).

297 We implemented two complementary techniques, regression coding schemes and
298 cutpoint structural analyses, to quantitatively examine temporal changes in somatic growth rates.

299 First, we used Reverse Helmert regression coding schemes to specifically compare growth rates
300 in the years before (1995–2009) and after (2010–2015) the *DWH* oil spill. The advantage of this
301 approach is that it allows for analysis of the entire growth dataset. We implemented coding
302 schemes using age class-specific linear mixed-effects models that included annual growth rate as
303 the dependent variable, year as the independent variable, and first-order autoregressive [AR(1)]
304 covariance structure for growth increments within turtles. Turtle-specific random effects were
305 also included to account for non-independence in the growth dataset—each turtle contributes
306 multiple growth rates. We then used maximally selected rank statistics to identify the optimal
307 cutpoint within each growth time series. This non-parametric approach was performed using the
308 mean growth rates for each age class, is robust to small sample sizes (Hothorn and Lausen 2003;
309 Müller and Hothorn 2004), and was implemented using the *coin* package in R (Zeileis et al.
310 2002; Hothorn et al. 2006).

311 Generalized Additive Models (GAMs) were used to examine relationships between
312 population density metrics and mean age class-specific growth rates. Models included age-
313 specific abundance (*Abund*) or cumulative hatchling production (*HatchProd*) as a fixed effect, an
314 identity link, and a quasi-likelihood error function. Within each model, mean growth rates were
315 weighted by sample size (i.e., number of growth rate estimates per year). For the oceanic life
316 stages (age 0, age 1), age-specific growth rates were compared to the model-derived cumulative
317 number of 0- and 1-year old turtles predicted to exist in a given year (metric 1) or the
318 cumulative number of hatchlings produced in a given year and the year prior (t_0-t_{-1}) (metric 2).
319 For the neritic life stages (age 2–5, age 6–9), age-specific growth rates were compared to the
320 model-derived cumulative number of juvenile turtles (ages 2–5) predicted to exist in a given

321 year (metric 1) or the cumulative number of hatchlings produced two to five years in the past ($t-$
322 $2-t-5$) (metric 2). Models were implemented in R using the *mgcv* package (Wood 2006).

323 We used cross-correlation to examine relationships between mean age class-specific
324 growth rates and climate indices. Following Bjorndal et al. (2016), GAMs with AR(1)
325 covariance structure were fit to the annualized climate data to reveal underlying trends since
326 1950 for the wNAO and AMO and since 1979 for the MEI. Mean age class-specific growth rates
327 were then compared to lagged (0- to 5-yrs) smoothing spline fits generated from the GAMs using
328 the *ccf* function in R (version 3.5.3; R Core Team 2019). Cross-correlation coefficients were
329 used to measure the degree of similarity between the two time series.

330 Lastly, to directly compare the independent and synergistic effects of these environmental
331 stressors on sea turtle growth rates, we performed an integrative analysis that incorporated the
332 results of the aforementioned independent analyses into a family of GAMs for each age class.
333 Models included various combinations of the three factors investigated as fixed effects, an
334 identity link, and a quasi-likelihood error function. We weighted mean growth rates by sample
335 size and used AIC and w_i to evaluate and compare models (Burnham and Anderson 2002).

336

337 **Results**

338 *Age and Growth*

339 SCL and age at stranding ranged from 4.2 to 69.1 cm SCL and 0 to 30.25 years for turtles
340 stranded on U.S. GoM beaches. Turtles stranded on U.S. Atlantic Coast beaches were 19.3 to
341 66.7 cm SCL and 1.00 to 18.75 years old (Tables 1, S1). Although their contribution to the
342 breeding population is not well understood (NMFS and USFWS 2015), documentation of tagged
343 Atlantic turtles nesting on the species' primary nesting beach in Mexico suggests that Atlantic

344 Kemp's ridleys ultimately return to the GoM as large juveniles or maturing adults (Caillouet et
345 al. 2015), resulting in relatively few adult animals on the Atlantic Coast. In total,
346 skeletochronological analyses yielded 3647 annual growth rate estimates from 1235 turtles for
347 the years 1988 to 2015 (Fig. 2). This constitutes the largest and most comprehensive dataset of
348 Kemp's ridley somatic growth rates to date. Annual growth rates span ages 0 (first year of life) to
349 28.75 but data from younger ages (< 6 yrs) dominate the dataset (~75%) because
350 younger/smaller turtles are the most abundant Kemp's ridley age classes in the population and
351 thus constitute the majority of stranded turtles (Gallaway et al. 2016).

352 For both the GoM and Atlantic Coast, there were distinct spatiotemporal changes in
353 humerus bone collection (Fig. 2). Prior to 2010, GoM samples were primarily obtained from
354 turtles stranded in Texas and Florida, whereas after 2010 sample collection shifted to turtles
355 stranded in Louisiana, Mississippi, and Alabama as part of the *DWH* oil spill response efforts.
356 Along the U.S. Atlantic Coast, there was a similar shift in sample collection in 2014 and 2015
357 from turtles that stranded primarily in North Carolina and Virginia to turtles that stranded in
358 Massachusetts. Using a general linear mixed model that accounted for year, age, AR(1)
359 autocorrelation, and turtle-specific random effects, we found somatic growth rates did not differ
360 within regions (Tukey's post hoc test, $p > 0.05$) but were significantly lower in turtles from the
361 Atlantic Coast (Tukey's post hoc test, $p < 0.05$). Examination of age class-specific growth rates
362 indicates that these regional differences in growth manifest as early as Age 1 and extend through
363 the small neritic juvenile life stage (age 2–5) (Fig. 3). Regional differences in Kemp's ridley
364 growth, size-at-age relationships, and maturation trajectories are further examined by Avens et
365 al. (*in review*), whose analysis uses the same growth rate dataset presented herein.

366 The quantity of age class-specific somatic growth rate data was sparse for years
367 preceding 1995, so all temporal growth analyses begin in 1995 and generally extend through
368 2014 or 2015 (Fig. 3). The datasets for age 0, age 2–5_{GoM}, age 2–5_{Atlantic}, and age 6–9_{GoM} turtles
369 are the largest and most continuous—all years have at least seven independent growth rate
370 estimates (Fig. 3). In contrast, significant data gaps exist for age 1_{GoM}, age 1_{Atlantic}, and age 6–
371 9_{Atlantic} turtles and the datasets for age 6–9_{GoM} and age 6–9_{Atlantic} turtles only extend to 2012 and
372 2010, respectively. We thus urge caution when interpreting results from the age 1 and age 6–9
373 datasets given that they are discontinuous and do not reflect similar time frames as the data for
374 ages 0 and 2–5.

375

376 *Deepwater Horizon oil spill effects*

377 The von Bertalanffy growth models fit to GoM turtle stranding length-at-age data
378 suggested that somatic growth differed before and after the *DWH* oil spill (Table 2, Fig. 4). The
379 model with the lowest AIC score and highest individual Akaike weight (w_i of 0.305) included
380 common L_∞ and t_0 parameters but different K parameters for the two time periods (1993–2009
381 vs. 2011–2016; Table 2). Parameter estimates for the best model were $L_\infty = 65.04$, $t_0 = 1.52$, K
382 (pre-*DWH*) = 0.192, and K (post-*DWH*) = 0.178. However, the next three best models had Δ AIC
383 scores less than 2.0 and w_i values between 0.113 and 0.298. While the parameters that differed or
384 agreed between the two time periods varied in these models (common L_∞ , different K and t_0 ;
385 different L_∞ , K , and t_0 ; common t_0 , different L_∞ and K), all included two separate K parameters.
386 The summed weights of the models that included separate K parameters for the two time periods
387 was 0.890, indicating overall support for a growth rate reduction in the GoM after the *DWH* oil
388 spill. Although we found some evidence for differences in K parameters, there was significant

389 overlap in the distributions of the stranding length-at-age data before and after the *DWH* oil spill
390 (Fig. 4), which suggests this apparent difference may not be biologically meaningful or that there
391 was not a systemic change in somatic growth across all U.S. GoM Kemp's ridley size classes.
392 Insufficient length-at-age data for larger/older Atlantic Kemp's ridleys, which are thought to
393 migrate back to the GoM prior to maturity (Caillouet et al. 2015), impeded our ability to fit von
394 Bertalanffy growth models for these turtles (but see Avens et al. *in review*).

395 Reverse Helmert regression coding schemes applied to the full somatic growth dataset
396 identified significant decreases in Kemp's ridley growth rates between 2011 and 2012 (Table 3).
397 Relative to pre-*DWH*, growth rates in 2012 declined by 1–2 cm yr⁻¹ within the age 0 and age 2–
398 5_{GoM} time series and greater than 3 cm yr⁻¹ within the age 2–5_{Atlantic} time series. Notably, this
399 analysis revealed that growth rates in 2013 (age 0, age 2–5_{GoM}) and 2014 (age 2–5_{GoM}, age 2–
400 5_{Atlantic}) were also significantly lower than pre-*DWH* growth rates. Relative to pre-*DWH*, growth
401 rates from 2012 to 2015 were lower by 8.1 % for age 0 turtles, 22.7 % for age 2–5_{GoM} turtles, and
402 30.7% for age 2–5_{Atlantic}. Similar results were obtained using complementary cutpoint analyses,
403 which identified significant decreases in mean annual somatic growth rates between 2011 and
404 2012 for turtles in the oceanic (age 0; max T = 3.14, p = 0.005) and small neritic juvenile life
405 stages in both the U.S. GoM (age 2–5_{GoM}; max T = 2.98, p = 0.008) and Atlantic Coast (age 2–
406 5_{Atlantic}; max T = 3.37, p = 0.004) (Table 3, Fig. 3). The cutpoint analysis did not identify any
407 statistically significant changes in somatic growth rates for the age 1 and age 6–9 time series (p <
408 0.05), though regression coding identified a significant increase in age 1_{Atlantic} growth rates and
409 decrease in age 6–9_{Atlantic} growth rates 2014.

410 Taken together, these analyses provide evidence for a sharp decline in Kemp's ridley
411 growth rates in the years following the *DWH* oil spill. However, the results of the temporal

412 analyses did not align with our original hypotheses that predicted either an acute (H_{1A}) or chronic
413 (H_{1B}) *DWH* oil spill impact on somatic growth rates beginning in 2010 for turtles in the GoM
414 only (Fig. 1). Interestingly, this decline is evident, and proportionally greater, in Atlantic
415 stranded turtles, which we predicted to exhibit no temporal changes in growth rates around the
416 time of the *DWH* oil spill due to their spatial isolation. However, even with a decrease in growth
417 rates, GoM small neritic juveniles (age 2–5) still grew faster than Atlantic conspecifics.

418

419 *Density-dependent effects*

420 We found little support for density dependent effects of cumulative turtle abundance and
421 hatchling production on mean age class-specific somatic growth rates (Table S2). For all but age
422 2–5_{GoM}, GAMs revealed no significant relationship between these population density metrics and
423 somatic growth ($p > 0.05$)—mean annual growth rates did not decline with increasing predicted
424 juvenile abundance nor was there the presence of a threshold above which growth rates declined.
425 The GAM response functions for both population abundance metrics and both GoM and Atlantic
426 stranded turtles were generally similar (Figs. S2–S4).

427 Cumulative hatchling production was a significant ($p = 0.018$) predictor of age 2–5_{GoM}
428 somatic growth whereas cumulative age 2–5 abundance was only a marginally significant ($p =$
429 0.051) predictor (Table S2, Fig. 5). Growth rates at the highest age 2–5_{GoM} population
430 abundances were lower on average than those at lowest predicted population abundance,
431 although 95% confidence intervals surrounding the annual means at the highest and lowest
432 abundances overlapped extensively. Nevertheless, the shapes of this relationship for age 2–5_{GoM}
433 did align with our hypothesis related to density-dependent effects (H₂) on somatic growth rates
434 (Fig. 1), which predicted a threshold above which growth rates begin to decline.

435

436 *Climate effects*

437 Mean annual growth rates tended to poorly correlate with the annualized climate indices
438 with 0- to 5-year lags (Table S3). Cross correlations for most life stages (age 1, age 2–5, age 6–
439 9) were generally negligible to weak (cross correlations $\leq |0.40|$), although cross correlations for
440 age 6–9_{GoM} with 4- and 5-yr lags were -0.53 and -0.59 for wNAO and 0.52 and 0.60 for AMO.
441 In contrast, mean annual growth rates exhibited moderate to strong correlations with all climate
442 indices for the oceanic life stage (age 0; Fig. 6). The highest, consistent cross correlation values
443 for age 0 included a 2-year lag (wNAO = 0.59 ; AMO = -0.57 ; MEI = 0.60). Cross correlations
444 values $\geq |0.50|$ were also observed for the wNAO and AMO with 3- to 4-yr lags, and the MEI
445 with 0- to 1-yr lags. The consistency in age class-specific growth patterns through time (Fig. 3)
446 generally do not align with our predicted climate growth response (Fig. 1: H₃) of declining
447 growth rates beginning in the late 1990s. However, our results suggest climate variability may
448 affect hatchling and oceanic juvenile growth during the oceanic life stage.

449 For the wNAO, positive cross correlations indicate that growth rates are higher when
450 winter weather conditions in the western North Atlantic are warmer and wetter (Drinkwater et al.
451 2003) and during periods of high river discharge, enhanced blue crab productivity, and reduced
452 *Sargassum* abundance in the GoM (Sanchez-Rubio et al. 2011, 2018). Similar conditions along
453 with cooler ocean temperatures are present during negative AMO phases (Karnauskas et al.
454 2015), which aligns with our observation of negative correlations between AMO and growth
455 rates (i.e., positive wNAO and negative AMO are coupled). Positive correlations between the
456 MEI and growth indicates growth rates increase with increasing ocean temperatures (Giannini et
457 al. 2001). However, our observation of declining oceanic stage turtle growth during a period of

458 warming suggest that indirect negative effects of increasing ocean temperatures on sea turtle
459 foraging habitat or prey may be negatively impacting their growth rates (Bjorndal et al. 2017).

460

461 *Integrative effects*

462 Three sets of GAMs were implemented to determine which environmental factors—
463 either independently or synergistically—were most strongly related to age 0, age 2–5_{GoM}, and
464 age 2–5_{Atlantic} growth. Comparative models were restricted to these age classes because they
465 showed evidence of significant temporal, density, and climate effects within independent
466 analyses. The three metrics evaluated in these models were (1) the temporal shift (*TS*) in growth
467 observed in 2012, included as a categorical variable ($TS_{pre} = 1995\text{--}2011$, $TS_{post} = 2012\text{--}2015$);
468 (2) cumulative hatchling production (*HatchProd*), included as a continuous variable; and, (3) the
469 annualized GAM trend for the AMO index with a 2-year lag, included as a continuous variable.
470 We generated models that included all combinations of these covariates as fixed effects, resulting
471 in the evaluation of six models for each age class (i.e., $TS + HatchProd + AMO$, $TS +$
472 $HatchProd$, $TS + AMO$, TS , $HatchProd$, AMO). The *HatchProd* and *AMO* covariates displayed a
473 moderate to high degree of collinearity with variance inflation factors of ~ 6 and correlation
474 coefficients between 0.83 and 0.92, indicating that the coefficients in the global model ($TS +$
475 $HatchProd + AMO$) may be poorly estimated and that the p-values may be questionable
476 (Dormann et al. 2013). We include the model herein for comparison but urge caution when
477 interpreting their results.

478 Within each age class, multiple models had ΔAIC scores less than two and were thus
479 considered strong potential predictors of mean age class-specific growth rates (Table 4). In all
480 cases these top models included *TS* alone or in combination with *HatchProd* and *AMO*.

481 However, in most cases *HatchProd* and *AMO* were not statistically significant predictor
482 variables (Table 5). The *HatchProd* and *AMO* only models explained the least variation in
483 somatic growth for all age classes.

484 For age 0, the top model included *TS* and *HatchProd* as fixed effects based on AIC score
485 and Akaike weight. However, the next three best models were within 2 AIC, which included *TS*
486 + *HatchProd*, *TS* only, or *TS* + *HatchProd* + *AMO* as fixed effects. The cumulative Akaike
487 weight for these top four models was 1.00. and *TS* was the only statistically significant predictor
488 of mean age 0 growth rates in these top four models (Table 5).

489 For age 2–5_{GoM}, the best model included all three covariates as fixed effects and had an
490 Akaike weight of 0.49. Notably, all three covariates were statistically significant predictors of
491 age 2–5_{GoM} somatic growth rates within this top model. A second model, *TS* + *HatchProd*, was
492 within 0.29 AIC of this best model. Only *TS* was a statistically significant predictor of mean age
493 2–5_{GoM} growth rates in this second model (Table 5), although *HatchProd* was marginally
494 significant ($p = 0.072$). The cumulative Akaike weight of these top two models was 0.92.

495 For age 2–5_{Atlantic}, the top model included *TS* only and had an Akaike weight of 0.45.
496 Two additional models had Δ AIC scores less than two (*TS* + *HatchProd* and *TS* + *AMO*),
497 providing for a cumulative Akaike weight of 0.86 for the top three models. As for the age 0
498 models, *TS* was the only statistically significant predictor of mean age 2–5_{Atlantic} growth rates
499 within the top age 2–5_{Atlantic} models (Table 5).

500

501 **Discussion**

502 Marine ecosystems are experiencing unprecedented change due to the combined effect of
503 suites of environmental factors. As population responses to ecosystem change are manifested

504 through changes in animal demographic rates, establishing mechanistic links between
505 environmental stressors and demographic variation is fundamental to understanding and
506 predicting species population dynamics. Through an analysis of 20+ years of somatic growth rate
507 data, we show that juvenile Kemp's ridley sea turtles experienced a significant, multi-year
508 reduction in somatic growth from 2012 to 2015 that spanned multiple life stages (oceanic and
509 small neritic juveniles) and habitats (GoM and U.S Atlantic). Specific mechanisms underpinning
510 this population-wide temporal shift in growth remain elusive, but likely include direct and
511 indirect negative effects of the *DWH* oil spill. Cumulatively, drivers of this 2012 change in
512 somatic growth constitute the single greatest contributor to somatic growth variation in recent
513 decades among the environmental factors investigated, though our integrative analysis indicated
514 that regional climate variability and changing population density have likely had synergistic
515 effects on oceanic (climate only) and small neritic (climate + population density) juvenile
516 somatic growth rates in the GoM. Our results contrast with other post-*DWH* oil spill studies that
517 observed immediate effects on growth rates in invertebrates and fish in 2010 but align with
518 observations of declining stranded turtle nutritional condition in the northern GoM beginning in
519 2012 (Stacy 2015), a phenomena of unknown origin but that would likely reduce growth rates.

520

521 *Growth and the Deepwater Horizon oil spill*

522 We hypothesize that the 2012 reductions in growth observed across the species' U.S.
523 range result partially from indirect negative effects of the *DWH* oil spill on sea turtle health
524 mediated by ecosystem changes. We initially predicted a direct *DWH*-associated growth
525 response would manifest in 2010 for GoM turtle life stages only given the coincidence of the oil
526 spill and annual initiation of sea turtle somatic growth, and the observation of immediate changes

527 in other species' demographic rates (e.g., Rozas et al. 2014; Brown-Peterson et al. 2016; Herdter
528 et al. 2017; Perez et al. 2017). However, the lack of a growth response in 2010 suggests the
529 *DWH* oil spill may not have had immediate, direct impacts on sea turtle growth rates. Still,
530 indirect negative effects are likely given the scale of the oil spill and whose impact may have
531 taken years to transcend food webs to influence sea turtle demographic rates.

532 Chronic exposure to *DWH*-associated environmental toxins may threaten the long-term
533 health of marine megafauna in the GoM, including sea turtles. Following the 1989 *Exxon Valdez*
534 oil spill, chronic exposure to weathered oil entrained in sediments delayed the recovery of a wide
535 range of taxa for decades due to long-term effects on species demographic rates (Peterson et al.
536 2003). Similar effects appear to be compromising the long-term health, reproductive success, and
537 survival of GoM bottlenose dolphins (Schwacke et al. 2014, 2017; Lane et al. 2015; Kellar et al.
538 2017). Much like other mobile marine predators, sea turtles were exposed to *DWH*-associated
539 environmental toxins for years following the oil spill due to its spatial overlap with key oceanic
540 and neritic foraging grounds that they continued to use (Shaver et al. 2013; Hart et al. 2014;
541 Wallace et al. 2017; Berenshtein et al. 2020). The leaching and resuspension of oil-contaminated
542 sediments represents a continued, long-term threat to coastal GoM food webs (Murawski et al.
543 2016; Rouhani et al. 2017; Romero et al. 2017). Additionally, both oceanic and neritic sea turtles
544 directly ingested spilled oil and absorbed polycyclic aromatic hydrocarbons (PAHs) into their
545 tissues (Ylitalo et al. 2017; Reich et al. 2017), which can cause adverse physiological effects in
546 animals such as reduced growth (e.g., Meador et al. 2006; Albers 2006).

547 Interestingly, the observed 2012 decline in somatic growth aligns with a simultaneous
548 deterioration of neritic stranded turtle nutritional condition and shift in sea turtle foraging
549 behavior in the northern GoM. Necropsies of juvenile Kemp's ridleys (25–60 cm SCL, ~2–9 yrs)

550 stranded in the northern GoM between 2010 and 2014 revealed significant reductions in the size
551 of turtle fat stores beginning in 2012 (Stacy 2015). Coincident with this change was a greater
552 association of sea turtles with fishing piers in Mississippi where turtles regularly attempted to eat
553 fishing bait (Coleman et al. 2016), a behavior previously linked to reduced growth rates in
554 Kemp's ridleys (Rudloe and Rudloe 2005). The integration of these findings with those herein
555 suggest a fundamental shift in the functioning of northern GoM food webs prior to 2012 that
556 impacted turtle foraging, nutritional condition, and inevitably growth. Causal factors for changes
557 in nutritional condition and foraging behaviors have not been identified, but the spatiotemporal
558 proximity to the *DWH* oil spill is conspicuous. Alternatively, these changes may relate to the
559 collapse of the Mississippi blue crab fishery in 2011, which has been attributed to freshwater
560 inundation from the opening of the Bonnet Carré Spillway (GSMFC 2015), but may also be
561 connected to negative effects of the *DWH* oil spill (Alloy et al. 2015; Giltz and Taylor 2017).
562 Comparisons of the nutritional status and growth histories of dead stranded turtles may improve
563 our understanding of temporal variability in Kemp's ridley growth dynamics.

564 Negative impacts of the *DWH* oil spill on oceanic habitats were severe and were
565 predicted to also impact the growth rates of oceanic stage turtles beginning in 2010. However,
566 much like the GoM small neritic juvenile life stage, we did not observe a significant decline in
567 oceanic stage turtle growth rates in 2010 but in 2012. All Kemp's ridleys associate with floating
568 *Sargassum* in GoM oceanic habitats for the first 1 to 3 years of life before recruiting to neritic
569 habitats along either the GoM or U.S. Atlantic Coast (Turtle Expert Working Group 2000; Avens
570 et al. *in review*). Following the oil spill, *Sargassum* tended to accumulate oil, become hypoxic,
571 and sink (Powers et al. 2013). The loss or compromise of this critical habitat would have
572 ultimately increased predation rates, reduced prey availability, and increased the energetic costs

573 of foraging during this life stage (Witherington 2002). Given the vulnerability of oceanic stage
574 turtles, the lack of a 2010 and 2011 growth response may indicate stronger initial *DWH* effects
575 on survival rather than growth (McDonald et al. 2017). Interestingly, *Sargassum* abundance was
576 anomalously high in 2011 and 2012 throughout the tropical North Atlantic, which should have
577 renewed these habitats and provided oceanic stage turtles with optimal conditions for growth and
578 survival (Witherington et al. 2012; Gower et al. 2013; Powers et al. 2013). That growth rates
579 instead declined in 2012 and 2013 suggests either lingering effects of the *DWH* oil spill on these
580 food webs or the influence of another environmental stressor (outlined below).

581 The observation of a strong, proportionally greater decline in Atlantic small neritic
582 juvenile growth in 2012 was unexpected given our initial assumption that their growth rates
583 would not change following the *DWH* oil spill. The specific causal factors for this decline remain
584 unknown but could be related to negative effects of the *DWH* oil spill on GoM *Sargassum*
585 habitats. It is well established that early nutrition can impact life-time growth through ‘silver
586 spoon’ effects (Larsson and Forslund 1991; Madsen and Shine 2000; McAdam and Boutin 2003;
587 Gaillard et al. 2003), and many of the Atlantic Kemp’s ridleys that exhibited reduced growth in
588 2012–2015 would have occupied GoM *Sargassum* habitats in 2010. Therefore, it is plausible that
589 cumulative impacts of the *DWH* oil spill on oceanic turtle habitats and physiology compromised
590 their long-term turtle health and were carried with them into non-impacted marine habitats both
591 within and outside the GoM (Putman et al. 2015). We still do not understand why Atlantic
592 Kemp’s ridleys grow more slowly than their GoM counterparts (reviewed in Avens et al. 2017,
593 *in review*), but the underlying cause could have interacted with carryover effects of the *DWH* oil
594 to amplify their cumulative growth response and cause the proportionally greater decline in

595 Atlantic Kemp's ridley growth rates. Improved understanding of drivers of Atlantic Kemp's
596 ridley growth variation will be key to disentangling potential effects of the *DWH* oil spill.

597

598 *Interactive effects of multiple environmental stressors*

599 Although the driver of the post-2012 shift in growth was identified as the greatest single
600 contributor to Kemp's ridley somatic growth variation over the past twenty years, we found
601 support for additive or synergistic effect of changing population density and climate variability
602 on GoM turtle growth rates. Indeed, our integrative analysis identified all three environmental
603 factors as significant predictors of GoM small neritic juvenile somatic growth. One hypothesis
604 for the recent fluctuations in Kemp's ridley nest counts after a period of exponential growth is
605 that the carrying capacity of the GoM has been reached for this species (Gallaway et al. 2016;
606 Caillouet et al. 2016, 2018). Empirical support for this hypothesis, however, has been lacking
607 due to insufficient data independent of the nesting trends, which are confounded after 2010 with
608 unknown effects of the *DWH* oil spill (but see Shaver et al. 2016). Within both independent and
609 integrative analyses we found strong support for a statistically significant relationship between
610 population density metrics and GoM small neritic juvenile growth. Specifically, we observed
611 lower, more variable growth rates at the highest population densities and a multi-year declining
612 growth trend that began in the mid-2000s, which generally aligns with our initial predictions and
613 observations in Avens et al. (2017). However, these findings are equivocal. Growth rates at the
614 highest population densities (2010–2015) overlap considerably with growth rates at the lowest
615 population densities (1995–1999) (Fig. 5). Therefore, more research is needed, especially
616 extensions of the skeletochronology dataset, before we can confidently assert Kemp's ridley
617 population density is influencing their somatic growth rates. Importantly, our findings contrast

618 with those that have suggested that density dependent processes have influenced this population
619 as early as the year 2000 (Caillouet et al. 2018; Caillouet 2019).

620 Climate variability may also influence both oceanic and small neritic juvenile Kemp's
621 ridley growth rates in the GoM, though our independent and integrative analyses provide
622 conflicting results. Recent studies have linked decades-long declines in sea turtle growth rates in
623 the Caribbean Basin to a late-1990s climate-driven ecological regime shift (Bjorndal et al. 2013,
624 2016, 2017). Herein, cross-correlations between lagged climate indices and somatic growth rates
625 identified moderate to strong correlations for oceanic stage turtles but climate was not a
626 significant predictor within the integrative analysis. The opposite pattern was observed for GoM
627 small neritic juveniles, where the climate indices were poorly correlated with somatic growth
628 within the independent analysis but identified as a significant predictor in the top integrative
629 model. Conflicting results for small neritic juveniles may be due in part to issues with
630 collinearity between population density and climate metrics in the top integrative model which
631 could inflate variance in model parameters for one or both variables (Dormann et al. 2012).
632 These issues aside, as ectotherms, sea turtle growth rates would generally be expected to
633 correlate with temperature-driven climate indices such as the AMO and MEI, particularly during
634 the oceanic stage when they occupy epipelagic habitats and have limited capacity to fight ocean
635 currents. Therefore, changes in growth rates for oceanic stage turtles may reflect the synergistic
636 effects of regional climate variability on oceanic habitats and lingering impacts of the *DWH* oil
637 spill, whereas changes in growth rates for GoM small benthic juveniles may have been more
638 strongly influenced by interactive effects of the *DWH* oil spill and population density.

639 Our analysis focused on three environmental stressors with wide-reaching influence, but
640 many other environmental factors likely contributed to Kemp's somatic variation during the

641 study period, particularly for Atlantic turtles. Anomalous heatwaves occurred in the western
642 North Atlantic in 2012 and 2016 that caused widespread ecosystem change, including shifts in
643 species distributions and recruitment (Mills et al. 2013; Pershing et al. 2015, 2018; Henderson et
644 al. 2017). Though effects of these heatwaves on sea turtles remain unknown, negative effects of
645 rising temperatures on Kemp’s ridley foraging habitats and prey could have indirectly impacted
646 their growth rates, a mechanism suggested to explain the declining growth trends in western
647 North Atlantic loggerhead, green, and hawksbill sea turtles in recent decades (Bjorndal et al.
648 2016, 2017). More broadly, local water temperatures hold the potential to substantially
649 contribute to somatic growth variation in this species given that its geographic distribution spans
650 greater than 20 degrees latitude (18°N to 43°N), thus a wide temperature range, and that parts of
651 the U.S. Atlantic Coast are warming faster than anywhere else in the world (Pershing et al.
652 2015). Establishing mechanistic links between sea turtle growth rates and local water
653 temperatures, such as through comparison of terminal humerus bone growth rates with local
654 temperature records, will be critical to understanding how sea turtles may respond to climate
655 change (Stubbs et al. 2020). A suite of other environmental factors—regional diet variation, prey
656 availability and distribution, intra-and inter-specific competition, genetics, migration distance
657 (see Avens et al. 2017, *in review*; Ramirez et al. 2020)—have also been identified as possible
658 contributors to Kemp’s ridley somatic growth variation and warrant further study.

659

660 *Implications of reduced somatic growth rates*

661 Whether the observed growth declines represent a biologically meaningful change
662 requires further evaluation. Somatic growth and body size influence a host of other demographic
663 processes, such as mortality rate, time to maturity, and fecundity, that cumulatively impact

664 individual fitness and species population dynamics (Madsen and Shine 2000; Dmitriew 2011).
665 Therefore, any alteration to an individual's growth trajectory has the potential to have cascading
666 effects on population demography. The growth rate declines we observed are well within the
667 natural variation for this species (reviewed in Avens et al. 2017) but their severity varied by life
668 stage. For example, oceanic stage turtle growth rates declined by ~8 % after 2012 but GoM and
669 Atlantic small neritic juvenile growth rates declined by ~20% and ~30%, respectively. Avens et
670 al. (*in review*) determined that the U.S. GoM vs. Atlantic Coast differences in somatic growth
671 may delay Atlantic Kemp's ridley maturity by 2 to 3 years relative to GoM counterparts. Herein,
672 post-2012 GoM Kemp's ridley growth rates are similar to those of pre-2012 Atlantic Kemp's
673 ridleys, which suggests a multi-year delay in maturation for GoM turtles is possible. Moreover,
674 the proportionally greater decline in Atlantic Kemp's ridley growth rates may further deepen
675 their life-long disadvantage relative to GoM conspecifics. Integration of somatic growth data into
676 demographic models may shed important light on the impacts of these growth changes on sea
677 turtle population dynamics and implications for conservation and management.

678

679 *Conclusion*

680 Through analyses of 30 years of dead stranded turtle humeri, we examined the somatic
681 growth response of the critically endangered Kemp's ridley sea turtle to multiple environmental
682 factors. We identified a simultaneous decrease in growth rates beginning in 2012 for oceanic and
683 small neritic juveniles that stranded in U.S. waters. We hypothesize that these changes are due in
684 part to deleterious effects of the *DWH* oil spill on sea turtles and their GoM habitats. For certain
685 life stages, this growth response may reflect synergistic effects of the *DWH* oil spill, climate
686 variability, and density-dependent processes. Our understanding of the links between the *DWH*

687 oil spill and sea turtle growth rates may be greatly enhanced through geochemical analyses (e.g.
688 PAHs, trace elements, isotopes) of turtle bone tissues, which may reveal direct evidence of
689 exposure to *DWH*-associated environmental toxins (e.g., Wise et al. 2014; Wilson et al. 2015;
690 López-Duarte et al. 2016; Romero et al. 2018). This study highlights the critical importance of
691 long-term, continuous collection of sea turtle humerus bones for status and threat assessment. To
692 date, the collection of dead stranded turtle humeri has been inconsistent across both space and
693 time and we lack any knowledge of growth rates of Kemp's ridleys within Mexican waters,
694 where approximately 18% of adult females and an unknown proportion of juveniles forage.
695 Within the U.S., widespread collection of Kemp's ridley bones ended in 2015 but resumption of
696 these efforts and initiation of bone collection in Mexico will be necessary to fully evaluate the
697 long-term influence of these environmental factors on sea turtle growth rates.

698

699 **Compliance with ethical standards**

700 *Conflict of interest.* The authors declare they have no conflict of interest.

701 *Ethical approval.* This article does not contain any studies with human participants or live
702 animals performed by any of the authors.

703

704 **Data availability**

705 The dataset generated and analyzed in the current study are available from the corresponding
706 author upon reasonable request.

707

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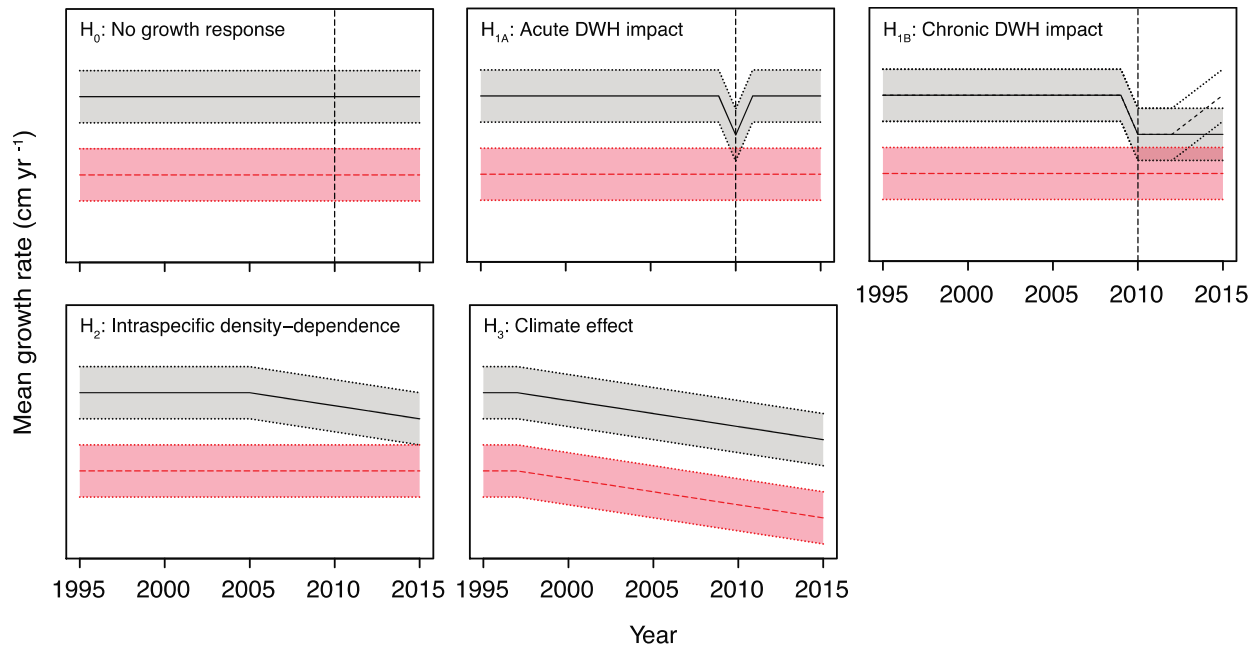
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Fig. 1 Conceptual model of alternative hypotheses for the size class-specific growth response of

1063

Kemp's ridley sea turtles to environmental factors examined herein. All Kemp's ridleys first

1064

reside in oceanic habitats in the central Gulf of Mexico (GoM) for 1–3 years then recruit to

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neritic habitats along either the GoM or U.S. Atlantic Coast. The shaded areas represent growth

1066

variation for GoM (black lines, grey shading) and Atlantic (red lines, red shading) life stages.

1067

Vertical dashed lines identify the year of the *Deepwater Horizon* (*DWH*) oil spill (2010). H_0 = no

1068

growth response in turtles from either geographic region or life stage to any factor examined. H_1

1069

= acute or chronic *DWH* oil spill-induced growth response for GoM life stages only (oceanic and

1070

neritic); no growth response in Atlantic neritic life stages due to geographic isolation from *DWH*

1071

oil spill, although Atlantic turtles may exhibit a past response during their GoM oceanic life

1072

stage. H_2 = density-dependent decline in somatic growth beginning in the mid-2000s during

1073

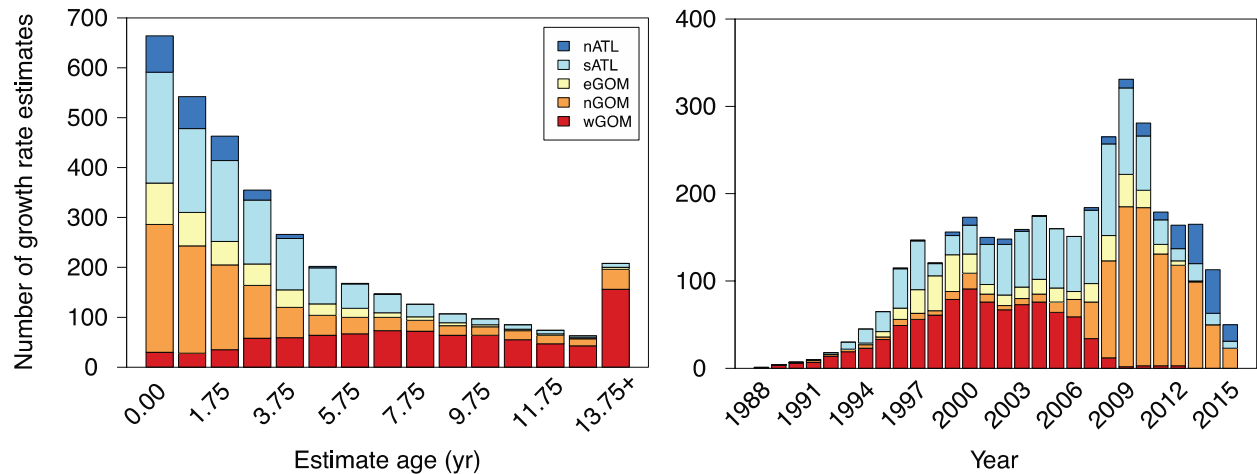
period of exponential population growth; effect in GoM turtles only as > 80 % of the population

1074

is thought to reside in the GoM (Putman et al. 2013; NMFS and USFWS 2015). H_3 = declining

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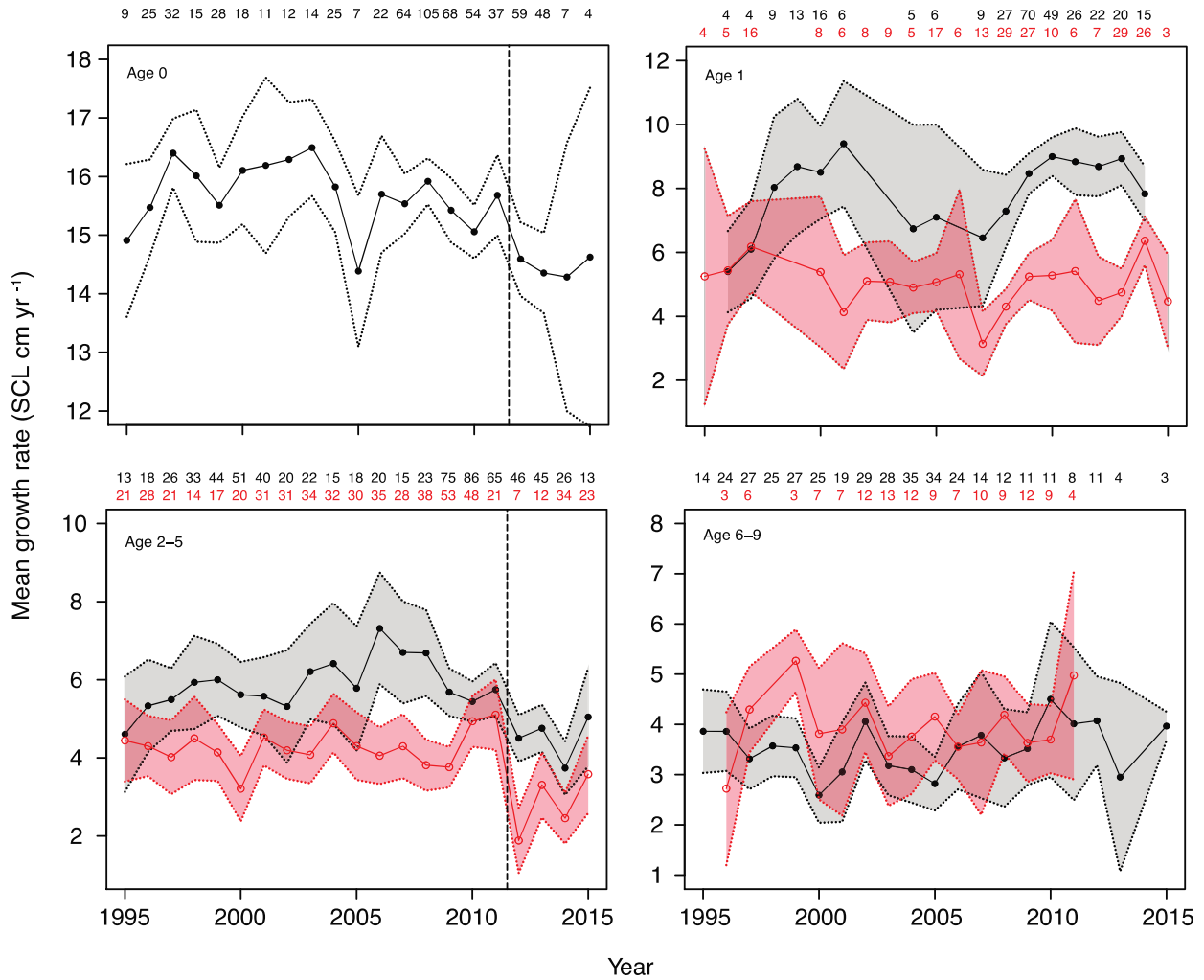
growth beginning in the late 1990s in response to climate-driven ecological regime shift



1076

1077 **Fig. 2** Frequency histograms of Kemp's ridley sea turtle back-calculated somatic growth rates by
 1078 stranding location, age, and year. nATL = northern Atlantic (stranding location = Delaware, New
 1079 Jersey, New York, Massachusetts), sATL = southern Atlantic (stranding location = Atlantic coast
 1080 of Florida, Georgia, South Carolina, North Carolina, Virginia), eGoM = eastern Gulf of Mexico
 1081 (stranding location = GoM coast of Florida), nGoM = northern Gulf of Mexico (stranding
 1082 location = Louisiana, Mississippi, Alabama), wGoM = western Gulf of Mexico (stranding
 1083 location = Texas)

1084



1085

1086 **Fig. 3** Time series of mean Kemp's ridley sea turtle growth rate by age class. Dotted lines bound

1087 95% confidence intervals. Age 0 includes data from both Gulf of Mexico (GoM) and Atlantic

1088 stranded turtles given that all Kemp's ridleys share oceanic habitats in the central GoM during

1089 the oceanic life stage. For all other age classes, GoM and Atlantic data were analyzed separately

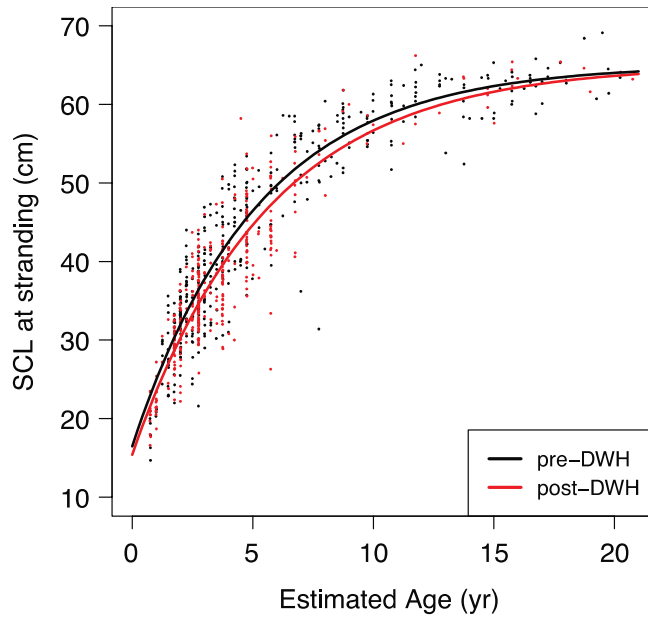
1090 due to regional differences in growth rates (black shaded area = Gulf of Mexico stranded turtles;

1091 red shaded area = Atlantic stranded turtles). Number of growth observations are presented above

1092 each plot. Vertical dashed lines identify significant breaks in each time series where there was

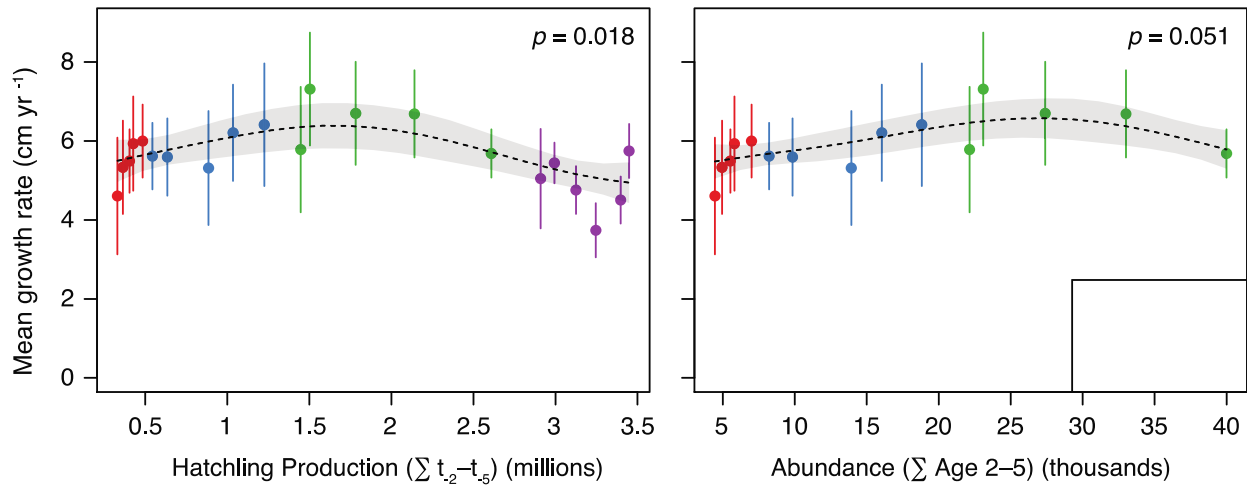
1093 concordance among statistical methods evaluated (see Table 3). Data for years with $N < 3$ are

1094 excluded. SCL = straightline carapace length

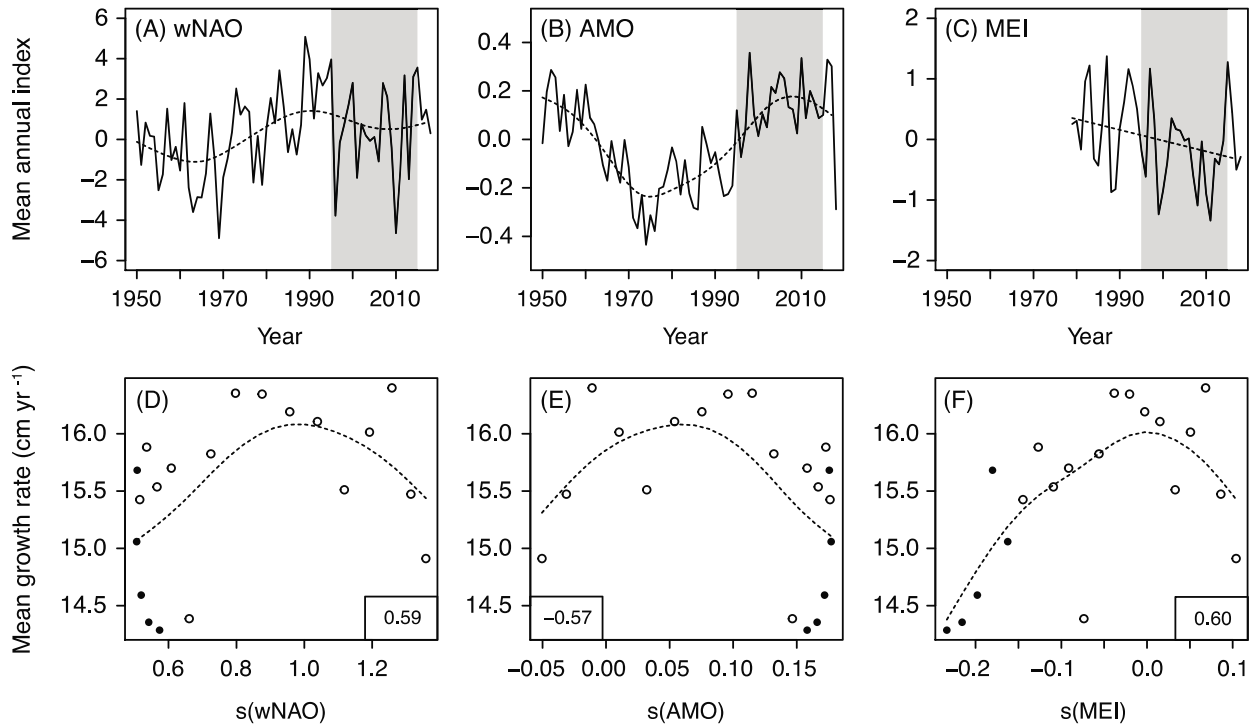


1095

1096 **Fig. 4** Von Bertalanffy growth functions estimated for Kemp's ridley sea turtles stranded in the
 1097 Gulf of Mexico before (1993–2009, $n = 402$) and after (2011–2016, $n = 362$) the *Deepwater*
 1098 *Horizon* oil spill. VBGFs were based on measured straightline carapace length (SCL) and
 1099 estimated age at stranding. Parameter estimates for the best model were $L_{\infty} = 65.04$, $t_0 = 1.52$, K
 1100 (pre-*DWH*) = 0.192, and K (post-*DWH*) = 0.178



1101
 1102 **Fig. 5** Relationship between mean back-calculated growth rate and population density metrics
 1103 for age 2–5 Kemp’s ridley sea turtles stranded in the Gulf of Mexico. Dashed lines and grey
 1104 ribbons are predicted values and 95% CI from GAM models with either cumulative hatchling
 1105 production (left panel) or population abundance (right panel) included as a smoother term (see
 1106 Table S2). Points are means ± 95% CI. SCL = straightline carapace length. See Figs. S2–S4 for
 1107 age 0, age 1, and age 6–9



1108

1109 **Fig. 6** Relationships between (A-C) climate indices and year and (D-F) mean age 0 growth rates

1110 and annualized climate indices (2-yr lag). Dashed lines are the GAM trends. (A-C) Shaded area

1111 identifies study period. (D-F) Cross-correlation values are presented in boxes within each plot.

1112 Open circles are years 1995–2009, whereas filled circles are year 2010–2015. wNAO: Winter

1113 North Atlantic Oscillation, AMO: Atlantic Multidecadal Oscillation, MEI: Multivariate El Niño

1114 Southern Oscillation Index.

Table 1 Summary characteristics for Kemp’s ridley sea turtles by stranding location. Western GoM (wGoM) = Texas; northern GoM (nGoM) = Louisiana, Mississippi, Alabama; eastern GoM (eGoM) = GoM coast of Florida); southern Atlantic (sATL) = Atlantic coast of Florida, Georgia, South Carolina, North Carolina, Virginia; northern Atlantic (nATL = Delaware, New Jersey, New York, Massachusetts). See Table S1 for state-specific data

Location	Stranding data				Growth rate data	
	<i>n</i> *	SCL (cm) Mean ± SD (range)	Estimated age (yr) Mean ± SD (range)	Year range	<i>n</i>	Year range
wGoM	200	55.6 ± 10.9 (4.2 – 69.1)	11.87 ± 6.47 (0.00 – 30.25)	1997 – 2013	915	1988 – 2012
nGoM	439	40.0 ± 11.1 (16.6 – 66.2)	4.86 ± 4.37 (0.75 – 23.00)	1993 – 2016	1055	1990 – 2015
eGoM	142	41.1 ± 11.0 (20.3 – 65.4)	4.62 ± 3.23 (1.00 – 15.75)	1998 – 2013	354	1994 – 2013
sATL	362	38.2 ± 10.3 (19.3 – 66.7)	5.07 ± 3.23 (1.00 – 18.75)	1993 – 2016	1071	1990 – 2015
nATL	77	28.0 ± 4.1 (19.3 – 40.0)	3.67 ± 1.41 (1.00 – 8.50)	2001 – 2017	219	1996 – 2015

*Stranding state unknown for 15 turtles (2 in Gulf of Mexico, 13 in Atlantic)

Table 2 Summary statistics for the family of models used to evaluate whether von Bertalanffy growth parameter estimates (L_∞ , K , t_0) differed for Kemp’s ridley sea turtles stranded in the Gulf of Mexico before (1993–2009, $n = 402$) and after (2011–2016, $n = 362$) the *Deepwater Horizon* oil spill. L_∞ is the asymptotic average length, K is the Brody growth rate coefficient, and t_0 is the age when the average length is zero.

Model	df	logLik	AIC	ΔAIC	W_i
Common L_∞ and t_0 ($K \neq K$)	5	-2201.34	4412.69	0.00	0.305
Common L_∞ ($K \neq K$, $t_0 \neq t_0$)	6	-2200.37	4412.74	0.05	0.298
Different L_∞ , K , and t_0	7	-2199.91	4413.82	1.13	0.174
Common t_0 ($L_\infty \neq L_\infty$, $K \neq K$)	6	-2201.33	4414.67	1.98	0.113
Common K and t_0 ($L_\infty \neq L_\infty$)	5	-2202.91	4415.81	3.12	0.064
Common K ($L_\infty \neq L_\infty$, $t_0 \neq t_0$)	6	-2202.50	4416.99	4.3	0.036
Common L_∞ and K ($t_0 \neq t_0$)	5	-2204.76	4419.52	6.83	0.010
Common L_∞ , K , and t_0	4	-2214.51	4437.02	24.33	0.000

Table 3 Results of Reverse Helmert regression coding schemes used to compare mean age class-specific growth rates of Kemp’s ridley sea turtles before and after the Deepwater Horizon oil spill. Number of asterisks (*) indicates degree of significance based on p-values (* = $p < 0.05$, ** = $p < 0.01$, *** = $p < 0.001$; empty cells mean no significant difference in mean growth rate). Colors indicate direction of change (**black** = increase, **red** = decrease). The complementary cutpoint analyses identified statistically significant structural shift in the age 0, age 2–5_{GoM}, and age 2–5_{Atlantic} growth time series between 2011 and 2012. Years without data for comparison with pre-*DWH* growth rates are noted with a dash

Year comparison	Age class						
	Gulf of Mexico stranded turtles				Atlantic stranded turtles		
Comparison	0	1	2–5	6–9	1	2–5	6–9
2005 vs. 1995–2004				*			
2006 vs. 1995–2005			**				
2007 vs. 1995–2006					**		
2008 vs. 1995–2007							
2009 vs. 1995–2008							
2010 vs. 1995–2009						***	
2011 vs. 1995–2009						**	
2012 vs. 1995–2009	**		*			**	
2013 vs. 1995–2009	***		**				
2014 vs. 1995–2009			***		**	***	*
2015 vs. 1995–2009							–

Table 4 Summary statistics for the family of Generalized Additive Models used to evaluate the influence of covariates [temporal shift (TS), hatchling production (HatchProd), Atlantic Multidecadal Oscillation (AMO)] on mean age class-specific growth rates for age 0 and age 2–5 Kemp’s ridley sea turtles. *TS* is a factor with categorization based on breakpoint identified in temporal analyses (pre-shift = 1995–2011, post-shift = 2012–2015). *HatchProd* is cumulative hatchling production for years, $t(x)$, prior to a given year (age 0 = $\sum t_0-t_{-1}$, age 2–5 = $\sum t_{-2}-t_{-5}$). *AMO* is the annualized GAM trend for the index with a 2-year lag.

Model	<i>df</i>	logLik	AIC	Δ AIC	W_i
(a) Age 0					
TS + HatchProd	4.00	–11.28	30.57	0.00	0.32
TS + AMO	5.34	–10.04	30.76	0.19	0.29
TS	3.00	–12.57	31.15	0.58	0.24
TS + AMO + HatchProd	5.00	–11.02	32.05	1.48	0.15
HatchProd	3.00	–20.72	47.45	16.88	0.00
AMO	3.39	–20.69	48.15	17.58	0.00
(b) Age 2–5, Gulf of Mexico					
TS + AMO + HatchProd	5.00	–11.17	32.35	0.00	0.49
TS + HatchProd	6.74	–9.58	32.64	0.29	0.43
TS	3.00	–15.51	37.03	4.68	0.05
TS + AMO	4.00	–14.85	37.71	5.36	0.03
HatchProd	5.46	–16.45	43.83	11.48	0.00
AMO	3.00	–23.51	53.02	20.67	0.00
(c) Age 2–5, Atlantic					
TS	3.00	–14.86	35.72	0.00	0.45
TS + HatchProd	4.00	–14.48	36.95	1.23	0.24
TS + AMO	4.00	–14.83	37.65	1.93	0.17
TS + AMO + HatchProd	5.00	–14.09	38.17	2.45	0.13
HatchProd	3.00	–21.41	48.82	13.10	0.00
AMO	3.00	–21.98	49.97	14.25	0.00

Table 5 Summary of statistical output for Generalized Additive Models (GAMs) used to evaluate the influence of potential environmental covariates [temporal shift (*TS*), hatchling production (*HatchProd*), Atlantic Multidecadal Oscillation (*AMO*)] on mean age class-specific growth rates for age 0 and age 2–5 Kemp’s ridley sea turtles. *TS* is a factor with categorization based on breakpoint identified in temporal analyses ($TS_{pre} = 1995–2011$, $TS_{post} = 2012–2015$). *HatchProd* is cumulative hatchling production for years, $t(x)$, prior to a given year (age 0 = $\sum t_0–t_{-1}$, age 2–5 = $\sum t_{-2}–t_{-5}$). *AMO* is the annualized GAM trend for the index with a 2-year lag. *Dev* : deviance explained by the model. *Edf*: estimated degrees of freedom. The models are ordered as in Table 4, with age class-specific models with ΔAIC scores < 2 denoted with an asterisk (*). Bold values denote statistically significant covariates ($p < 0.05$)

Model	Dev (%)	Adj. R^2	Smooth terms				Parametric coefficients					
			Var	Edf	F	Prob(F)	Var	Est	SE	t	Pr> t	
(a) Age 0 ($n = 21$ years)												
GAM _{TS} + HatchProd*	65.1	0.61	HatchProd	1.00	2.35	0.142	TS _{pre}	1.13	0.22	5.12	<0.001	
GAM _{TS} + AMO*	69.0	0.64	AMO	1.90	2.34	0.235	TS _{pre}	1.13	0.22	5.26	<0.001	
GAM _{TS} *	60.6	0.59	–	–	–	–	TS _{pre}	1.20	0.22	5.40	<0.001	
GAM _{TS} + AMO + HatchProd*	66.0	0.60	AMO	1.00	0.43	0.522	TS _{pre}	1.15	0.23	5.08	<0.001	
			HatchProd	1.00	1.30	0.270						
GAM _{HatchProd}	14.3	0.10	HatchProd	1.00	3.18	0.091	–	–	–	–	–	
GAM _{AMO}	14.6	0.09	AMO	1.21	1.68	0.165	–	–	–	–	–	
(b) Age 2–5, Gulf of Mexico ($n = 21$ years)												
GAM _{TS} + AMO + HatchProd*	69.2	0.64	AMO	1.00	8.69	0.009	TS _{pre}	0.99	0.29	3.38	0.004	
			HatchProd	1.00	7.14	0.016						
GAM _{TS} + HatchProd*	73.6	0.67	HatchProd	3.07	2.53	0.072	TS _{pre}	1.07	0.28	3.84	0.001	
GAM _{TS}	53.5	0.51	–	–	–	–	TS _{pre}	1.29	0.28	4.67	<0.001	
GAM _{TS} + AMO	56.3	0.52	AMO	1.00	1.17	0.294	TS _{pre}	1.39	0.29	4.80	<0.001	
GAM _{HatchProd}	49.1	0.41	HatchProd	2.84	4.11	0.018	–	–	–	–	–	
GAM _{AMO}	0.4	–0.05	AMO	1.00	0.07	0.793	–	–	–	–	–	
(c) Age 2–5, Atlantic ($n = 21$ years)												
GAM _{TS} *	50.0	0.47	–	–	–	–	TS _{pre}	1.39	0.32	4.36	<0.001	
GAM _{TS} + HatchProd*	51.8	0.46	HatchProd	1.00	0.68	0.422	TS _{pre}	1.55	0.38	4.10	<0.001	
GAM _{TS} + AMO*	50.1	0.45	AMO	1.00	0.06	0.808	TS _{pre}	1.40	0.34	4.19	<0.001	
GAM _{TS} + AMO + HatchProd	53.5	0.45	AMO	1.00	0.64	0.433	TS _{pre}	1.69	0.42	4.03	<0.001	
			HatchProd	1.00	1.24	0.280						
GAM _{HatchProd}	6.7	0.02	HatchProd	1.00	1.36	0.257	–	–	–	–	–	
GAM _{AMO}	1.4	–0.04	AMO	1.00	0.27	0.604	–	–	–	–	–	