Foraging Fidelity as a Recipe for a Long Life: Foraging Strategy and Longevity in Male Southern Elephant Seals

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Abstract

Identifying individual factors affecting life-span has long been of interest for biologists and demographers: how do some individuals manage to dodge the “little devils” of death [3] longer than the large majority of their conspecifics? To start answering this question, several problems need to be overcome; the first being the accurate estimation of life-span or longevity. In the case of vertebrates, longevity may be estimated by following individuals from their birth till their death using mark-recapture methods [4]. Then we used stable isotopic measurements of carbon (δ13C) in dentin to study the effect of foraging location on individual life-span. Using a joint change-point/survival modelling approach which enabled us to describe the ontogenetic trajectory of foraging, we unveiled how a stable foraging strategy developed early in life positively covaried with longevity in male Southern Elephant Seals. Coupled with an appropriate statistical analysis, stable isotopes have the potential to tackle ecological questions of long standing interest but whose answer has been hampered by logistic constraints.

Introduction

Identifying individual factors affecting life-span has long been of interest for biologists and demographers [1,2]: how do some individuals manage to dodge the “little devils” of death [3] longer than the large majority of their conspecifics? To start answering this question, several problems need to be overcome; the first being the accurate estimation of life-span or longevity. In the case of vertebrates, longevity may be estimated by following individuals from their birth till their death using mark-recapture methods [4]. However, long-lived organisms (for example, seabirds) present additional challenges: a wandering albatross (Diomedea exulans) may live up to 60 years [3], requiring a lot of patience, serendipity, and skills to secure funds and manpower from demographers. Yet knowledge of individual longevity may be critical to shed light on life-history patterns [6–12]. Recent studies have moved away from population-level (life-tables, for example [13]) to individual-level inferences (for example, [14]), which is the level where natural selection occurs. This move is the result of both conceptual and technical advances, most notably in estimating a notoriously difficult individual fitness [15,16]. It also results from the availability of rich datasets collected on wild populations over several decades. Such data depth allows to study the evolution and the ecological correlates of life-history traits in the wild.

In the case of mammal species, most studies investigating the relationship between longevity and fitness have focused on females [6–8,10–12,17–19] (but see [9]). These studies, which mainly concerned large terrestrial herbivores, usually found evidence of long-lived females having a larger fitness than short lived ones (but see [7]). On the other hand, males are usually not studied as estimating their fitness is harder and often demands genetic analysis to reliably infer offspring’s paternity.

The Southern Elephant Seal (Mirounga leonina) is the most dimorphic and polygynous mammal among extant species. The biology of this elusive carnivore, which can spend up more than 80% of its lifetime at sea [20], means that seals are not observable most of the time. Moreover, when ashore during the breeding season, males fight to hold harems of numerous females. Most males never reproduce but a few mates with a large number of females [21]. Body size is a critical component for holding and fighting over a harem. Since these seals can grow all their life [22], breeding for a male depends on surviving long enough to reach an adequate size to be able to hold a harem. We may therefore expect a strong relationship between longevity and fitness in males [23]. Assessing longevity in the male Southern Elephant Seal using mark-recapture methods is extremely demanding: less than 4% of a cohort may survive up to 9 years-old [24], when they may become harem-holders [25]. Moreover the question of why these males manage to outlive the others is left open since their at-sea behaviour remains elusive.

One way to overcome this problem is to rely on indirect methods to infer the at-sea ecology of these animals. In this
respect, the study of marine mammals has greatly benefited from the use of stable isotopes [26]. Carbon \(^{13}C\) and nitrogen \(^{15}N\) stable isotope ratio are the most commonly used elements in isotopic dietary studies. Carbon (nitrogen) stable isotopes can provide information about the diet’s geographical origin (trophic position) of a consumer [27]. In the Southern Ocean, the existence of a latitudinal gradient in carbon stable isotopes across water masses [28–30] allows researchers to infer where elephants seals have been foraging prior to hauling out [31]. The temporal window reflected in stable isotope values depends on the sampled tissue [32]. Therefore, questions that may be addressed with stable isotopes are tied to the careful choice of an appropriate tissue.

Stable isotopic measurement of tissues that are metabolically inert after synthesis, such as teeth or baleen, can yield information on the ecology of marine mammals over their whole life [33–37]. Incremental tissues of teeth may further allow age estimation [38–40]. Measuring stable isotopes in teeth may permit to investigate the ecological correlates of longevity. In the present work, we studied the influence of foraging strategy, as inferred from carbon stable isotopes measured in tooth, on the individual lifespan of male Southern Elephant Seal breeding on ıles Kerguelen, Southern Ocean. Bailleul et al. [31] found that males from ıles Kerguelen were mainly foraging either in Subantarctic waters (mostly the Kerguelen Plateau) or on the Antarctic Plateau (see Figure S1). In a previous study [41], we investigated the ontogeny of foraging behaviour. Here we investigated whether the observed dual strategy affects the longevity of males Southern Elephant Seals.

**Results**

**Growth Mixture Modelling**

Bailleul et al. [31] investigated the foraging behaviour of juveniles males using both remote-sensing tags and blood carbon isotopes. Blood \(^{13}C\) values for young small-sized males had a unimodal distribution while there was evidence of a bimodal distribution for older large-sized males. Dentin \(^{13}C\) values had a unimodal distribution for individuals younger than 3 years old, and a clear bimodal distribution after that age (Figures 1 and 2). However, this approach only described the growth pattern of \(^{13}C\) conditional on a foraging strategy. Furthermore it made the restrictive assumption that the growth curve shape of individuals with the same foraging strategy is identical. However this model was used for descriptive purposes, and below we present the results of an explanatory approach which aimed at identifying ecological correlates of longevity.

**Joint Modelling**

We adopted a joint modelling approach for analyzing lifespan [42–47]. We used a hierarchical change-point model [48,49] to describe individual time-series of \(^{13}C\) values (see Figure 3A), which enabled us to identify an ontogenetic shift between a juvenile stage and an adult one when males were committed to either an Antarctic or a Subantarctic strategy [41]. For the adult stage, the regression slope of \(^{13}C\) values against age is either close to zero, which means a very stable strategy of foraging in either Subantarctic or Antarctic waters, or negative reflecting the preponderance of foraging in Antarctic waters (see Figure 3A).

With an AFT, survival times are directly modelled, which eases the interpretations of coefficients; but a parametric distribution family must be specified [50,51] in contrast to the semi-parametric Cox Proportional Hazard (PH) model [52]. We nevertheless opted for the AFT model and assumed survival times to follow a Weibull distribution [43], whose core assumption is a monotonic hazard function [53], which seems reasonable for these data (Figure 4B and Figure S2). We embedded the AFT within a hierarchical change-point model for \(^{13}C\) values [41] (see the annotated code in Supplementary Materials).

The best survival model was the joint change-point/survival model (Table 1), but the model fit as assessed using Kolmogorov-Smirnov test was poor (\(D = 0.57, p < 0.001\), Figure S3). Further investigations also revealed modest to strong correlations (\(>0.6\)) between the \(\beta_k\) and \(\theta_k\) (Figure S4). Close inspection of the individual-specific parameters \(a_{k},\theta_{k}\) revealed that the only parameter to truly covary with longevity was \(\theta_{k}\), the slope after the ontogenetic shift (Figure 4B). This was further checked and confirmed with Stochastic Search Variable Selection [54] (not shown). Hence only the posterior estimate of \(\theta_{k}\) is reported in Table 2. Since \(\theta_{k}\) was negative, males with a very stable foraging strategy \((a_{k} \approx 0)\) had on average a larger life-span than the other males.

**Discussion**

**Ecological correlates of longevity**

As expected, male Southern Elephant Seals showed a clear mixture of two foraging strategies as they aged [31,55]. Using carbon stable isotope measurement from tooth, we found that some males had an Antarctic signature \((\approx -20: -19\%)\), while others had a Subantarctic signature \((\approx -17: -16\%)\). The pattern in Figure 1B suggested that \(^{13}C\) values reflective of an Antarctic signature increased in proportion over time. Such a pattern may reflect the progressive disappearance of males foraging elsewhere than in Antarctic waters, either because of an ontogenetic shift in foraging behaviour [56] or because of differential survival of males with different foraging strategies. To investigate this matter further, we adopted a joint change-point/survival modelling approach to explicitly relate the age of an individual [39] with a proxy of its foraging behaviour [34,35].

Males foraging in Antarctic waters didn’t have a longer longevity than males foraging in the Subantarctic waters. The increasing proportion of Antarctic \(^{13}C\) values (Figure 1) was more the result of small sample size [57] for advanced age classes and of ignoring within-individual correlation (recall that a seal of a given age can contribute up to 4 isotopic values because of the sampling design, see **Materials & Methods**). A joint modelling approach, which accounted for the longitudinal nature of our data, revealed a relationship between the stability of a foraging strategy and longevity. Seals that exhibited little variation in their tooth \(^{13}C\) values (see Figure 3A) were the most long-lived. A change-point model evidenced a negative correlation between the age at ontogenetic shift and the slope after this shift [41]: seals that had an early shift were constant in their foraging behaviour for the rest of their lives. Thus, this sophisticated modelling approach confirmed what an “eye-ball” analysis suggested: profiles with the smallest isotopic variation were from seals with the longest life-span (Figure 1A).

The two modes that progressively appear with age on Figure 1B reflects how seals that became faithful to a foraging strategy early in life lived longer that others. Thus, the two modes in the distribution of \(^{13}C\) values (Figure 1B) partly arose from the selective disappearance of males with a variable \(^{13}C\) profile.

This pattern of an early shift in life associated with far-reaching consequences in later-life underscore how crucial are the first years.
of life in this species [58]. There was in fact a small (in magnitude) positive correlation between the (positive) slope before and the (negative) slope after the shift [41]. The positive slope before the shift was expected because weaned pups rely exclusively on maternal milk before weaning. Phocid milk is very rich in lipids [59,60], which are depleted in the heavier carbon isotope [61]. Thus the positive slope before the shift may in part reflect the progressive independence from maternal resources [34,40,62,63]. Pups which became early on independent from maternal resources, were able to forage on their own and adopted a very stable foraging strategy. Those same pups also lived longer than the others, suggesting thereby the potential importance of early life history on latter performances [64–66].

Bradshaw et al. [67] studied the fidelity of adult female Southern Elephant Seals to their foraging grounds. Using a measure of overlap between visited zones along at least two consecutive foraging journeys, Bradshaw et al. [67] directly evaluated how females seemed to behave according to simple navigation rules, that is how females were “rational” in the sense that their behaviour was predictable. One major finding of this elegant analysis was the lack of a relationship between mass gain and spatial overlap between two successive foraging trips: females showed fidelity to a foraging ground irrespective of foraging success. Although Bradshaw et al. [67] lacked data on long-term survival and lifetime reproductive success of these females, which were of the same age-period-cohort to limit potential confounders, they speculated that the stability shown by these females may have arisen in their early life. While our study is confined to males, it is in agreement with the results of Bradshaw et al. [67]. Despite a cruder spatial resolution compared to tracking data, stable isotopes enabled us to look into the ontogeny of foraging strategy in male Southern Elephant Seals, and thus to evidence how foraging fidelity was associated with longevity.

We were nonetheless surprised that the Antarctic strategy was not associated with an increased life-span. At least for females, an Antarctic strategy may yield higher fitness pay-offs. There is a latitudinal gradient in pup weaning mass with pups born in colonies closer to Antarctica having a larger weaning mass on average than pups born at lower latitude rookeries [68]. As weaning mass correlates with first-year survival [69], this suggests

![Figure 1](https://example.com/figure1.png)

**Figure 1. Spaghetti and density plots of the tooth \( \delta^{13}C \) values from male Southern Elephant Seals.** The distribution is unimodal up to age 3 but changes to a bimodal distribution afterwards. Observations belonging to the component with the smallest mean are first in a minority but progressively increase in proportion until becoming the majority.
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that resources in Antarctic waters may be more profitable. Chaigne et al. [70] analyzed blood stable isotopes of juvenile Southern Elephant Seal males. Their study design was cross-sectional, but Chaigne et al. [70] showed that older males, as assessed from their body length, were more likely to forage in Antarctic waters compared to younger ones. They have interpreted this pattern as an ontogenetic shift in foraging grounds, which is consistent with the stable isotope analysis of dentin [41]. Foraging in Antarctic waters thus seems to be favored by bigger males, possibly because of higher fitness pay-offs.

Yet our longevity data did not suggest an increased survival of Antarctic foragers. Unlike females which remained in the marginal sea-ice zone, juvenile Southern Elephant Seal males from ıˆles Kerguelen readily foraged in the pack ice [55]. Getting trapped in the ice is a potential cause of mortality that males foraging over the Kerguelen Plateau or at the Polar Front do not face [31]. Our present analysis does not point to different foraging grounds influencing male life-span. The pattern uncovered is one of the benefit of a very stable foraging strategy with no deviation from an early age in life. This pattern may lend support to a spatial familiarity hypothesis [71,72], although a direct experimental test of such an hypothesis is currently not possible with Southern Elephant Seals.

That isotopic profiles covaried with longevity thus suggests that variability in foraging strategy is costly. Such costs may arise from unfamiliarity with novels environments, such as a greater susceptibility to predators, or increased travel costs. A non-exclusive alternative is that some individuals were more able to extract resources efficiently from the environment, either in Antarctic or Subantarctic waters. None of these interpretations suppose a strategy to be superior to the other in terms of fitness return. However, the second interpretation implies that seals which are less efficient to acquire resources may switch between foraging strategies while those which are efficient have no reason

\[ \delta^{13}C \]

**Figure 2. Mean \( \delta^{13}C \) values of the two-components growth mixture in relation to Southern Elephant Seal age.** Posterior means along with 95% Highest Probability Density (HPD) intervals are depicted. For advanced age classes (>9 years), this model suggested an increase in mean \( \delta^{13}C \) values for Subantarctic foragers, and a decrease for Antarctic foragers. This effect seems artefactual in light of Figure 1 where isotopic values are stable after age 9. The artefact results from the restrictive assumption on the growth curve shape (see Methods).

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to do so. Under this latter interpretation, the δ13C profile of seals may reflect their ‘quality’, quality being understood as a static trait that positively correlated with fitness [73]. We chose the term quality over fitness here as the reproductive success of males in our sample is unknown. Yet given the breeding biology of male Southern Elephant Seals, living long enough to grow large and defend a harem is a pre-requisite to contribute offspring to the next generation [23].

Limitations

A potential confounding factor in our data is that teeth were sampled from dead animal on beaches, thus we had to assume our sample was representative of the larger population of all males on îles Kerguelen. Assuming otherwise would imply that male found dead on beaches were different than those dying at sea. The average longevity in our sample was 7 years, and only 5 males were older than 10 years (Figure S5). In their study on reproduction costs on Sea Lion Island (52°26’S, 59°05’W), Galimberti et al. [25] found only 4 males out of 78 (~5%) to be older than 10 years old, while McCann [74] reported a proportion of ~22% for South Georgia (54°15’S, 37°05’W). The observed proportion in our sample was 11(4,24)% , compatible with both the Sea Lion Island and South Georgia estimates. Age in our study was estimated from teeth growth layers: there is an uncertainty associated with age (~1 year). It is, however, very small [40] and cannot reverse the observed pattern. Defining a species’ longevity as the time by which 99% of a cohort has died [75], the specific longevity of male Southern Elephant Seals is 13 years [76]. The oldest male in our sample was estimated to be 12 years old, which suggested that our sample did not seem atypical with respect to old age classes.

Of concern may be the lack of fit of the AFT model to the data. Even our best model in term of AICc did not provide an adequate fit to the data (see Figure S3). Yet it has been argued that the poor predictive ability is an intrinsic feature of survival models with realistic parameter values [77]. Our approach here was explanatory rather than predictive [78]. The joint model clearly captured some aspect of the data unaccounted for by the Null model given its large Akaike weight. Further model checking revealed that this model was overparametrized, but still performed better than the null model despite the penalty for the larger number of (unnecessary) parameters. The mixture model also has a larger likelihood than the null model, but its larger number of parameters put it on a par with the null model (similar Akaike weights). Thus
the data suggest an effect of the foraging strategy that we seemed to have picked up best with a change-point model.

**Conclusion**

Stable isotopes, while lacking the fine scale resolution of tracking data, can reveal surprising ecological features of a species [26]. However, this crudeness may prove a strength: by summarizing a whole foraging trip with a single number, isotopic data provided an integrative measure that can be easily fed into a model specifically tailored to the problem at hand. Using an appropriate tissue, stable isotopes can also provide longitudinal data [33–37].

The explicitly modelling of foraging strategy ontogeny in male Southern Elephant Seal via a change-point (or broken-stick) model of δ13C values revealed how long-lived animals were those faithful to a foraging strategy from an early age. This finding emphasizes the importance of early life in life-history trajectories. It also suggests that variability in foraging strategies might be costly for adult male Southern Elephant Seals from îles Kerguelen. In other words, faithfulness to a foraging strategy predicted a long life for males.

Studying of the life-span of wild animals is a difficult enterprise: ecological correlates can be uncovered but a large amount of variation usually remains unaccounted for in the analysis [79]. This is unsurprising in light of all the potential factors, related to fitness or accidental, that may affect an individual throughout its whole life [3,12,80].

**Materials and Methods**

**Ethics Statement**

The ethics committee of the French Polar Institute (Institut Paul Emile Victor - IPEV) approved this study. All animals in this study were cared for in accordance with its guidelines. This study is part of a national research program (no. 109, H. Weimerskirch and the...
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Table 1. Accelerated Failure Time (AFT) model selection.

<table>
<thead>
<tr>
<th>Model</th>
<th>K</th>
<th>Deviance</th>
<th>AICc</th>
<th>ΔAICc</th>
<th>wAICc</th>
</tr>
</thead>
<tbody>
<tr>
<td>Joint</td>
<td>6</td>
<td>202.2</td>
<td>215.6</td>
<td>0.0</td>
<td>94.5</td>
</tr>
<tr>
<td>Null</td>
<td>2</td>
<td>218.4</td>
<td>222.5</td>
<td>6.9</td>
<td>3.0</td>
</tr>
<tr>
<td>Mixture</td>
<td>4</td>
<td>215.6</td>
<td>224.1</td>
<td>9.0</td>
<td>1.4</td>
</tr>
<tr>
<td>Random</td>
<td>3</td>
<td>218.3</td>
<td>224.6</td>
<td>9.5</td>
<td>1.1</td>
</tr>
</tbody>
</table>

Four models were considered and we kept a ratio of number of parameters to be estimated to the number of datum close to 8. The Akaike Information Criterion corrected for small sample size (AICc) is reported. ΔAICc is model AICc minus the minimum observed AICc. wAICc are model weights. The best model in terms of predictive ability was the joint change-point/survival model.

As a recent study raised concerns about non-linear offsets of natural causes on îles Kerguelen (49° 30′ S, 69° 30′ E), Southern Indian Ocean. Canines grow continuously throughout the whole life of males without closing of the pulp cavity, allowing for age determination [81]. Canines from 47 males were analyzed and sampled for isotopic analysis [see [40] for a full description of age determination [81]. Canines from 47 males were analyzed and sampled for isotopic analysis [see [40] for a full description of age determination and isotopic sampling]. Briefly, each tooth was cut longitudinally and observed under diffused light to count growth layers.

Sample Collection

Teeth were collected from male elephant seals that died of natural causes on îles Kerguelen (49° 30′ S, 69° 30′ E), Southern Indian Ocean. Canines grow continuously throughout the whole life of males without closing of the pulp cavity, allowing for age determination [81]. Canines from 47 males were analyzed and sampled for isotopic analysis (see [40] for a full description of age determination and isotopic sampling). Briefly, each tooth was cut longitudinally and observed under diffused light to count growth layers. The alternating pattern of two opaque and two translucent growth layers corresponds to the annual biological cycle of Southern Elephant Seals [39]. Translucent bands are enriched in vitamin D and synthesized when seals are ashore to breed and to molt, while opaque ones are synthesized when at sea [82].

Within a year, a Southern Elephant Seal comes onshore to molt, while opaque ones are synthesized when at sea [82]. Thus each growth layer was assumed to correspond to one forth of a year [40]. Each growth layer was sampled for 1 mg of bulk dentin using a MicromillTM sampler (ISEM, Université de Montpellier 2).

As a recent study raised concerns about non-linear offsets of organic %C, %N and C/N after acid treatment [83], we forwent any acid (or demineralization) treatment prior to isotopic measurement. As a result, the measured 13C is a mixture of organic carbon with a small amount of inorganic carbon. To test the impact of the inorganic fraction, Martin et al. [40] compared acid-treated and untreated samples but found no differences (± 0.02). Schulting et al. [84] found similar C/N ratios between bulk dentin and collagen, with a lower carbon and nitrogen contents in bulk dentin most likely due to the mineral fraction. Here we assumed that the impact of the mineral fraction is negligible.

For measurement of carbon stable isotopes and C/N ratio, a total of 1,233 dentin increments over the 47 male teeth were analyzed. Elemental C and N contents (%) and carbon isotope values were measured by dry combustion using a Euro Vector 3000 Elemental Analyzer coupled with a Micromass Optima Ratio Mass Spectrometre (ISEM, Université de Montpellier 2). Results are expressed in percentage of powder weight (Total C and N) and as 13C (‰) with respect to the Vienna-Pee Dee Belemnite standard using the conventional delta notation:

\[ \delta^{13}C = 1000 \times \left( \frac{R_{\text{sample}}}{R_{\text{standard}}} - 1 \right) \]

where \( R_{\text{sample}} \) and \( R_{\text{standard}} \) refer to the 13C/12C ratios of sample and standard, respectively. Analytical precision was better than 0.20‰.

We used C/N ratio thresholds of bone and tooth collagen (2.9 to 3.6) as criteria for the identification of diagenetic alteration [83]; assuming that total dentin, whose organic phase is mainly collagen and water [86], has the same C/N ratio than bone and tooth collagen. From the 1,233 analyzed samples, 118 were discarded, yielding a final sample size of 1,115 isotopic values from 47 males. Given the alternating pattern of tooth growth layers, up to 4 isotopic measurements were available for a given year of life.

Table 2. Summary statistics for the joint change-point/survival model.

<table>
<thead>
<tr>
<th>Parameters</th>
<th>Median</th>
<th>2.5%</th>
<th>97.5%</th>
<th>Interpretation</th>
</tr>
</thead>
<tbody>
<tr>
<td>( \beta )</td>
<td>3.3</td>
<td>2.6</td>
<td>4.2</td>
<td>Shape of the Weibull distribution</td>
</tr>
<tr>
<td>( \beta_0 )</td>
<td>-6.7</td>
<td>-8.6</td>
<td>-5.0</td>
<td>Intercept</td>
</tr>
<tr>
<td>( \beta_4 )</td>
<td>-1.4</td>
<td>-2.8</td>
<td>-0.1</td>
<td>Slope after shift</td>
</tr>
</tbody>
</table>

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Figure S1). We thus assumed a two-components mixture. For the $j^{th}$ male, we modelled its blood isotopic signature at age $t$ as:

$$\delta^{13}C_{jt} = A_{jt} + z_{jt} \times e^{t \lambda_{jt}} + \epsilon_{jt}$$

where

- $\epsilon_{jt}$ is a normally distributed residual term with 0 mean and variance $\sigma^2_{res}$
- $A_{jt}$ is the isotopic value at birth
- $A_{jt}$ is the logarithm of the absolute isotopic difference between 12 and 0 years
- $z_{jt}$ are weights describing the growth curve for individuals foraging in Subantarctic waters
- $z_{jt}$ are weights describing the growth curve for individuals foraging in Antarctic waters
- $k$ is a latent indicator variable of the foraging strategy

For model identification purposes, we constrained $z_{0,k=1} = 0$, $z_{1,k=1} = 1$ and $z_{2,k=1} = -1$. In addition, to circumvent any label switching issues, we further constrained the isotopic value of Antarctic foragers to be lower than that of Subantarctic foragers. Practically, we allowed the batch of weight coefficients of the first component of the mixture to be positive ($0 \leq z_{jt} \leq 1$) but forced the second batch to be negative ($-1 \leq z_{jt} \leq 0$). Finally, we assumed the growth curves to be isotonic: for all age $t$, $z_{jt} \leq z_{jt+1,t}$ and $z_{jt} \geq z_{jt+1,t}$. Isotonicity translates an assumption about isotopic equilibration to a foraging habitat signature. A crucial assumption of this approach is that the growth curve shape of individuals with the same foraging strategy is identical, which is not reasonable given the variety of profiles observed in Figure 1. To relax this assumption, we used a hierarchical random change-point model [41].

**Joint Modelling**

Change-point models aim at finding an abrupt rupture in a time-series. The time-series is assumed to be the juxtaposition of piece-wise linear homogeneous segments, each segment separated from the next by a change-point. These models are very flexible as they allow specifying different probability distributions to describe different parts of a time series. Different curve shapes can thus be generated. Change-point models thus seem especially appropriate to describe ontogenetic shifts [41,56]. A time-series is summarized in 4 parameters: a value at the change-point, the timing of the change-point, and a slope before and after the change-point. In a previous paper [41], we used a hierarchical change-point model to describe individual time-series of $^{13}$C measurement in Southern Elephant Seal teeth and found evidence of ontogenetic shifts. Here, we assessed the impact of these ontogenetic shifts on longevity.

Because teeth were sampled from dead animals that were subsequently aged, all survival times are observed: there is no censoring in the data. Denoting $T_j$ the survival time of the $j^{th}$ male, we assumed the $T_j$ to follow a Weibull distribution of parameters $r$ and $\lambda_j$:

$$T_j \sim \text{Weibull}(r, \lambda_j)$$

$$\log(t_j) = \beta_0 + \beta_1 \times a_{jt1} + \beta_2 \times a_{jt2} + \beta_3 \times a_{jt3} + \beta_4 \times a_{jt4}$$

where

- $a_{jt1}$ is isotopic value at change point
- $a_{jt2}$ is slope before the change point
- $a_{jt3}$ is logarithm of the age at change point
- $a_{jt4}$ is slope after the change point

The shape parameter $r$ controls the hazard rate with $r > 0$ (resp. $r < 0$) describing an increasing (resp. decreasing) hazard with time. With our data, we expected $r > 0$ (Figure 4). The parameters $\beta_1, \beta_2, \beta_3, \beta_4$ then quantify the association between foraging location (via $\delta^{13}$C) and longevity. The parameter $\beta_4$ captures the relation between age at ontogenetic shift and longevity. The parameter we are particularly interested in is $\beta_4$ as it reflects the correlation between the stable foraging habitat of adults and their longevity. In the AFT, a positive $\beta \times x$, where $x$ is the covariate value, accelerates the occurrence of the event (death), while a negative value retards it. A negative $\beta_4$ means that individuals foraging in two different water masses (negative slope) die earlier than those males which have a very stable strategy (null slope). The joint modelling approach is especially suited here as it makes use of all the available data and the Bayesian framework guarantees that uncertainties in estimating the $a_{jt1,2,4}$ are taken into account [43,46].

To assess the appropriateness of our joint model, we compared it with 3 other AFT survival models:

- • a null model with no individual-level covariate;
- • a random-effect model wherein an individual-specific deviation from the mean $\delta^{13}$C value was incorporated as a covariate for the AFT model; and lastly
- • a mixture model wherein each male was first assigned to a group depending on its mean $\delta^{13}$C value and then group membership was included as a covariate for the AFT model.

Strictly speaking, both the random and mixture models are also joint models, as time-series of $\delta^{13}$C values were used to derived predictors for the survival analysis. Model comparison were done using the Akaike Information Criterion with a small sample correction, AICc [88]. Our sample size is modest ($N=47$) and the most complex AFT model considered had 6 parameters, keeping the ratio of sample size to parameter number above $\approx 8$ which is slightly below the recommended 10 [53]. The goodness-of-fit of the selected model was checked by comparing the predicted longevity with the observed one using Kolmogorov-Smirnov test. Finally, we investigated in a preliminary analysis whether males born before and after the 1970s population crash [89] had different longevity and found none (Likelihood Ratio Test: $\chi^2 = 0.30, p = 0.86$).

Growth layer synthesized while seals were ashore were kept in all analyses. These layers may differ from the others since Southern Elephant Seals fast on land. Retaining these layers may add measurement error linked to physiological processes. We then compared for each seal the distribution of isotopic values measured in dentin synthesized ashore versus at-sea with a Kolmogorov-Smirnov test. Except for one individual, there was no statistically significant differences (See Table S1). Excluding this individual did not change our results. Balasse et al. [90] estimated the isotopic equilibration of dentin after diet change to take between 1 to 4 months in cattle (Bos taurus). This time-span is commensurate with, if not longer than, the typical haul-out duration of a Southern Elephant Seal. Assuming similar equilibration time for cattle and elephant seals, this may explain
why no statistically significant differences were found (See Table S1).

Software

We used winBUGS [91] called from R [92] with the package R2WinBUGS [93]. Weakly informative priors were used [94,95]. For the growth mixture model, we used a uniform prior for the residual variance; Normal priors for regression parameters on the natural scale; the default Student-t prior of [95] for regression parameters on a logarithmic scale; and a beta(2,2) prior for the mixing proportion. We used the SVD prior of Tokuda et al. [96] for the covariance matrix controlling the $\Lambda_j$ random orthogonal matrices were generated as described in Anderson et al. [97]. Three chains were initialized with overdispersed starting values. After appropriate burn-in (40,000 iterations) and thinning of the chains (1 value every 10 iterations stored), convergence was assessed using the Gelman-Rubin convergence diagnostic [98] with the coda package [99]. For the joint model, Authier et al. [41] detailed the hierarchical change-point model fitted to the isotopic data. We used for the AFT model the default Student-t prior of [95] for the parameter $\beta_{d+s+r}$ and a uniform prior bounded between 0 and 10 for r. Three chains were initialized with overdispersed starting values. After appropriate burn-in (200,000 iterations) and thinning of the chains (1 value every 200 iterations stored), convergence was assessed using the Gelman-Rubin convergence diagnostic [98] with the coda package [99]. Unless stated otherwise, posterior mean and standard error of the mean are reported, either with its standard error ($\pm$ se) or with 95% Highest Probability Density (HPD) intervals (Mean$^{\pm2.5\%,97.5\%}$). Inferences are based on a posterior sample of 3,000 iterations. Annotated BUGS code is available in Text S1 (Growth Mixture Models) and Text S2 (Joint Change-point/Survival Model).

Supporting Information

Figure S1 Satellite tracking of Southern Elephant Seal males breeding on îles Kerguelen. Examples of 24 tracks are represented (solid blue lines) to illustrate the different strategies: males mainly forage in the Antarctic Zone, on the Kerguelen Plateau or in Subantarctic waters (waters lying between the Sub-Antarctic Front and the Southern Antarctic Circum-Polar Front); îles Kerguelen (Ker), and the Antarctic coastline’s contour are depicted in black, and while the 2000 meters isobath is depicted in light grey. Dotted lines symbolized fronts [103], withi n the Southern Ocean: SubTrop-ical Front (STF), Sub-Antarctic Front (SAF), Polar Front (PF) and Southern Antarctic Circum-Polar Front (SACCF). Margin- al histograms of localizations are represented on the side to illustrate the different strategies.

Text S1 WinBUGS code to fit the Growth Mixture Model. (XLS)

Text S2 WinBUGS code to fit the joint Accelerated Failure Time\Change-point Model. (XLS)

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Author Contributions

Conceived and designed the experiments: IB CG. Performed the experiments: IB CM AF. Analyzed the data: MA. Contributed reagents/ materials/analysis tools: CM IB CG. Wrote the paper: MA IB CG.

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