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Author for correspondence:

James C. Bull

e-mail: j.c.bull@swansea.ac.uk

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THE ROYAL SOCIETY

Climate causes shifts in grey seal phenology by modifying age structure

James C. Bull¹, Owen R. Jones^{2,3}, Luca Börger¹, Novella Franconi¹, Roma Banga¹, Kate Lock⁴ and Thomas B. Stringell⁵

(D) JCB, 0000-0002-4373-6830

There are numerous examples of phenological shifts that are recognized both as indicators of climate change and drivers of ecosystem change. A pressing challenge is to understand the causal mechanisms by which climate affects phenology. We combined annual population census data and individual longitudinal data (1992-2018) on grey seals, Halicheorus grypus, to quantify the relationship between pupping season phenology and sea surface temperature. A temperature increase of 2°C was associated with a pupping season advance of approximately seven days at the population level. However, we found that maternal age, rather than sea temperature, accounted for changes in pupping date by individuals. Warmer years were associated with an older average age of mothers, allowing us to explain phenological observations in terms of a changing population age structure. Finally, we developed a matrix population model to test whether our observations were consistent with changes to the stable age distribution. This could not fully account for observed phenological shift, strongly suggesting transient modification of population age structure, for example owing to immigration. We demonstrate a novel mechanism for phenological shifts under climate change in long-lived, age- or stage-structured species with broad implications for dynamics and resilience, as well as population management.

1. Introduction

There are numerous examples of changes to the timing of biological events from wide-ranging taxonomic groups and it is well understood that these shifts can provide sensitive indicators of the effects of climate change, e.g. [1–4]. The causes and consequences of phenological shifts resulting from climate change have become a major area of interest in recent years, across many ecosystems and geographical regions; for examples of recent reviews and syntheses, see [5–9]. Now, a pressing challenge is to develop a robust, mechanistic understanding of how these processes operate [10–12]. This is needed to predict changes in abundance and distribution and to facilitate conservation of endangered or otherwise important species [13].

Additionally, species do not exist in isolation and phenological changes can cascade through biological communities, via trophic, competitive, and mutualistic interactions [14–16]. In particular, mismatches in seasonal events, for example between predator and prey populations or flowering plants and their pollinators, can decouple biological communities and lead to critical transitions in population structure, biological regime shifts, and even collapse of ecosystem services [17–19]. In ecological communities that are strongly regulated by top-down processes, climate change-driven shifts in predator distributions and dynamics are likely to have downstream effects on whole assemblages of species and ecosystem

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¹Department of Biosciences, Swansea University, Swansea SA2 8PP, UK

²Department of Biology, University of Southern Denmark, Odense, Denmark

³Interdisciplinary Centre on Population Dynamics (CPOP), University of Southern Denmark, Odense, Denmark

⁴Natural Resources Wales, Martin's Haven, Pembrokeshire, UK

⁵Natural Resources Wales, Bangor, Gwynedd, UK

functions [10,14,20–22]. Therefore, understanding the mechanistic drivers of phenological shifts in key predatory species is of fundamental importance for our ability to meet the global challenge of conserving biodiversity and ecosystem function under climate change.

Here, we focus on grey seals (Halichoerus grypus), as a wellstudied example of marine mammals, e.g. [23-29]. Like many other seal species, grey seals are iteroparous capital breeders with a high degree of site fidelity [30]. They start breeding at around 5 years old and reproduce annually for potentially several decades, resulting in populations with overlapping generations [31,32]. Marine mammal phenology has been cited as evidence of a major system shift in the Indian Ocean and parts of the Southern Ocean [33] and climate-driven changes in seasonal timing are predicted to have substantial impacts on marine mammal populations themselves [32,34]. There are now observable changes in the timing of seal life history across the Atlantic [35-37], including phenological shifts associated with climate [32], making this an important focal group in their own right, as well as important indicators of the wider effects of climate change.

Phenological shifts in life-history events, including breeding [37–39], pupping [40] and moulting [41] have been reported in many seal species. Many species of seal are known to undergo delayed implantation, or embryonic diapause [42–44], in common with other large carnivore species [45–49]. This has been proposed as the mechanism by which climate acts on pupping phenology in seal species [26]. It is hypothesized that colder sea temperatures invoke a physiological response which delays implantation, thus the mode of action operates at the individual seal level [26]. However, whether temperature alone is sufficient is not certain [49,50], and it is hard to test whether this potential mechanism is responsible for observed population level phenological shifts in highly mobile and elusive species.

An alternative hypothesis has been proposed: that the mechanism of pupping season phenology acts at the population level, through climate-driven modification of the local population age structure, rather than at the individual level, e.g. [26,35]. While the effect of climate change on age structure has been shown to underpin phenological shifts in birds [51,52], and fishes [53], the age structure hypothesis has rarely been tested in mammals, including seals, and not supported where it has [35]. If found to occur in nature, this would open an important new line of enquiry into the mechanisms driving phenological responses to climate change in long-lived predators with age- or stage-structured life histories.

A possible reason why the age structure hypothesis has not been supported in seals previously is a focus on local populations and demographic processes [35]. In closed populations, describing the stable age structure may be enough to understand dynamics, brought about through local demographic processes, such as changes to fecundity and differential survival across age classes. However, along with many age- or stage-structured species, grey seals are known to roam widely [54-56] and even where site fidelity is high, local populations are open to immigration and emigration as a further source of changes to age structure [30]. In grey seals, younger adults have been found to be more likely to remain resident and have smaller home ranges than older adults [57]. This suggests a mechanism where older animals are more likely to move to favourable sites as environmental conditions change, increasing the mean age of the population at those locations. Coupled with older grey seals having higher weaning success [31], and observations that older individuals tend to reproduce earlier across many taxa, including seals [58–62], it seems reasonable to hypothesize that climate change may act on phenology and population dynamics through modification of age structure, underpinned by movement patterns.

The aims of the current study are to: (i) quantify how changes in the timing and progression of the grey seal pupping season are dependent on climatic drivers throughout a long-term monitoring programme; (ii) identify whether phenology acts at the individual level or population level in a natural setting; and (iii) test the hypothesis that modification of population age structure is the causal mechanism for observed phenological changes. While we do not have data to test whether it is local demography or movement that changes age structure, we focus our modelling on whether stable age structure can account for our findings and discuss the likely role of movement in explaining that.

2. Methods

(a) Study site

The Skomer Marine Conservation Zone (SMCZ) area in Pembrokeshire, south Wales, UK (51°43′55.2″ N, 5°16′33.6″ W) comprises the island of Skomer and mainland Marloes Peninsula (figure 1) and is the location of one of the largest grey seal pupping sites in Wales [54]. Adult female grey seals haul out on sheltered beaches and caves throughout this area to give birth and nurse pups until weaning after about three weeks [55]. The pupping season around Pembrokeshire runs from late August to December (figure 1, inset calendar).

(b) Seal data

Each year, trained staff carry out surveys most days during the pupping season (typically every 1–3 days from late summer to the end of the year). For this study, records of seal pup counts were used for the period 1992–2018 (27 years) (electronic supplementary material, figure S1). Individual records were kept of each pup's progress from birth to moult (approx. three weeks) following a standardized protocol, e.g. [54]. Details included a unique numerical pup identifier, the location (beach name) and date of first observation, the developmental stage (from 1 to 5) of the pup [56], date of subsequent observations, and a record of neonatal survival until weaning.

On Skomer Island, seal pups were marked with a unique pattern using coloured aerosol sheep fleece marker sprays to help monitor each individual. Additionally, adult seals were identified using their distinctive pelage patterns, which persist as unique identifiers throughout adult life, particularly on females [63]. This approach allowed multi-annual records of individual adult female pupping timing to be compiled. On the Marloes Peninsula, pups are spread across a series of small, inaccessible beaches, surrounded by cliffs. Pup numbers on each beach are low enough for individuals to be recognized by experienced staff on subsequent days without marking. This allows accurate recordings to be made during a given season but follow-up of adults between years was not formalized for Marloes seals [55]. Seal pup analysis for this study was conducted across the whole SMCZ, while analysis of adults in this study was conducted on Skomer seals only. Approximately 72% of the seal pup productivity in this study was on Skomer Island.

Some wider context about SMCZ seals is available as electronic supplementary material, Study species.

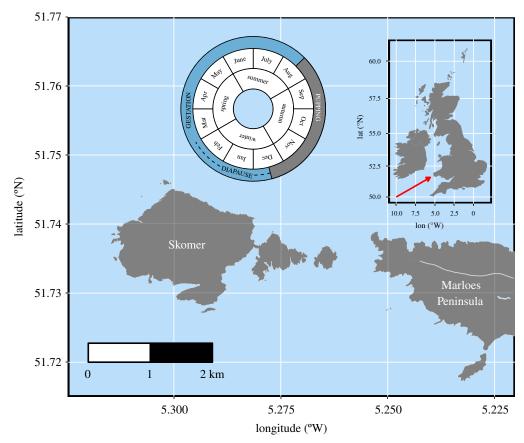


Figure 1. Skomer Island and the Marloes Peninsula, southwest Pembrokeshire, Wales. Inset map shows the location of the study area within the British Isles (red arrow). Inset circular plot shows the pupping season for grey seals around Pembrokeshire relative to our defined seasons: 'autumn' = September–November (the majority of the pupping season), 'winter' = December–February, 'spring' = March–May and 'summer' = June–August. (Online version in colour.)

(c) Environmental data

HadISST1 satellite measurements of sea surface temperatures (SST) over the period of this study were obtained from the British Atmospheric Data Centre (http://www.ceda.ac.uk/data-centres/). These data are resolved into geographical cells (1° latitude \times 1° longitude). We used the cell with the northwest corner (52° N, 6° W), covering the whole of the SMCZ. SST data (electronic supplementary material, figure S2) were aggregated into quarterly averages, anchored around the height of the pupping season: 'autumn' = September–November (pupping season), 'winter' = December–February, 'spring' = March–May and 'summer' = June–August.

(d) Statistical analysis

Observations of seal pupping were used to analyse the cumulative increase in pup counts throughout the pupping season for each year. All dates were described as days since 1 July in a given year. However, because the end of the observing season was dictated more by logistical constraints than the date of the last pup birth, we developed a model to estimate the progression of the pupping season that was insensitive to random fluctuations in the tails of the season.

Empirical cumulative counts of pups were seen to closely follow a sigmoid shape each year (electronic supplementary material, figure S3). Therefore, we fitted three-parameter logistic curves to cumulative pup counts, separately for each year, using nonlinear mixed-effects models with 'year' as a random variable [64]. We accounted for within-year serial dependency as a first-order autoregressive error process, and between-year heteroscedasticity by estimating within-year variances separately. The three parameters that describe these curves are the asymptotic total number of pups, the point of inflection of the fitted sigmoid

curve, and a scale parameter regulating the steepness of increase of the curve. The point of inflection provides a robust estimate of the midpoint of the season (the date where 50% of pups have been counted) and the scale parameter is proportional to the length of the season. We assessed goodness of fit using concordance correlation coefficients [65], with greater than 99.5% concordance in all years.

Changes in each of these three yearly derived parameters (pupping season total, midpoint and length) were modelled separately using generalized additive models (GAMs). We fitted SST as an explanatory variable to test our hypothesis and included the estimate of pupping season total from the previous year as an additional explanatory variable, to account for autocorrelation [37], potentially owing to a number of factors, including autocorrelated environments, energetic carry-overs, or unaccounted for species interactions. Mating occurs after pupping, which could introduce a lagged effect on season midpoint and length. Therefore, we also included the season midpoint and season length estimates from the previous year in our statistical models of season midpoint and length, respectively. All explanatory variables were modelled using cubic smoothing splines. In all cases, we modelled residual error distributions using gamma distributions with natural logarithm link functions.

Here, stepwise model selection was inappropriate, as models are not nested owing to differences in the degree of nonlinearity [66]. In such cases, shrinkage smoothers are recommended [67], which were used here. Separate models were constructed for each of the quarterly SST estimates ('previous autumn', 'winter', 'spring', 'summer' and 'autumn' = current pupping season) and compared using Akaike information criterion (AICc). Since 'winter' was found to result in the lowest AICc, we proceeded using this season. This is also consistent with the hypothesis that the three months following mating (which occurs in the autumn, after pupping) determines the subsequent pupping date [26].

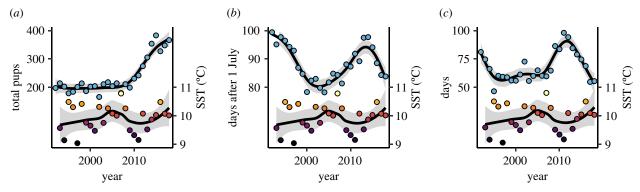


Figure 2. Grey seal pupping season parameters and sea surface temperature (SST) around the Skomer Marine Conservation Zone, from 1992 to 2018. Left-hand vertical axes show season parameters: (a) the total pups over the season, (b) the midpoint of the season (days since 1 July) and (c) an estimate of 95% of the season length (days). Blue data points show season parameter estimates. Right-hand vertical scale shows winter SST (December–February). The fill colour of temperature data points (bottom of plots) scales with SST. Solid lines showing fitted trends and shaded ribbons representing marginal 95% confidence intervals. (Online version in colour.)

To assess population-level demographic responses to climate, we used longitudinal data on re-sightings of individual adult females. Here, 'age' is measured as years since first sighting. We modelled how the average age of adult females around Skomer varied with SST using a GAM. We fitted natural logarithm-transformed mean adult age as the response variable, with winter SST as an explanatory variable using cubic smoothing splines. We modelled residual error using Gaussian distributions with identity link functions.

In cases where individual adults were observed to give birth over several years, we were also able to assess whether the date at which an individual gives birth changes with age, or in response to SST, as well as the statistical interaction between ageing and SST. We modelled the date of pupping by individual adults around Skomer using generalized additive mixed models (GAMMs) with gamma error distributions and natural logarithm link functions. We fitted years since the first sighting of a given individual adult and SST as additive fixed effects using cubic smoothing splines. As the age of adults at first sighting is unknown, we also incorporated individual adult female seal identity as a random effect. This allowed us to describe their response to fixed effects without assuming a common starting age (intercept).

(e) Age-structured population modelling

To explore the relationship between demography and the phenology of the breeding season, we constructed a seven-stage matrix population model (MPM) (electronic supplementary material, table S1). This model serves as a useful tool to explore the potential effects of altered vital rates on population structure, and consequently on population-level breeding phenology. The model is age-based and includes four juvenile stages and three adult stages. The first juvenile stage represents weaned age (approx. 3–4 weeks old) to 1 year, while the others represent years. Reproduction from the three adult stages varies, accounting for the probability of successful weaning depending on maternal age. Therefore, our model's transitions from these adult age-classes to the 'weaning-1 year' stage include both the probability of giving birth and neonatal survival until weaning.

Full details of MPM parameterization are available as the electronic supplementary material, Matrix population model parameterization.

The age structure of the seal population could influence the population-level mean pupping date if individual-level pupping date is associated with maternal age. Therefore, we conducted a sensitivity analysis of this MPM to explore how changes in vital rates, and consequently age structure (stable age distribution), might influence population-level average pupping date. We calculated the expected population-level average pupping date by using

the stable age distribution as weights in a calculation of mean pupping date given our knowledge of age-specific pupping dates.

Specifically, we calculated the change in population-level mean pupping date resulting from a +10% increase in each matrix element. For transitions with a negative sensitivity (i.e. that could result in an earlier pupping date), we then asked how much earlier could the pupping date potentially become? We did this by calculating the effect on pupping date of increasing these survival transitions up to the maximum reasonable value of 0.999.

All statistical analysis was undertaken using R v. 3.6.0 [68] with additional packages: AICcmodavg v. 2.3–1 for model comparisons and selection [69], mgcv v. 1.8–31 for GAM(M)s [70], ncdf4 v. 1.17 for reading SST data from .nc files [71], and popdemo v. 1.3–0 for matrix population modelling [72].

3. Results

(a) Phenological trends

Overall, the SMCZ grey seal pupping season showed marked variation throughout the study period, 1992-2018 (figure 2). The annual total grey seal pup counts showed a substantial increase in the last quarter of the survey period, averaging around 200 pups until 2009, then doubling by 2015 (figure 2a). The estimated midpoint of the season was in the first week of October at the beginning of the survey (latest date: 7 October in 1992), advancing by about three weeks to mid-September through the middle part of the survey (earliest date: 16 September in 2004), and returning to early October towards the end of the survey (figure 2b). The estimated pupping season length showed a twofold change throughout the monitoring period: 47 days in 1995, 98 days in 2011 (figure 2c). For all three phenological parameters, the respective parameter value from the previous year had a statistically significant effect (total pup estimate: F = 51.7, p < 0.001; season length: F = 11.4, p < 0.001; season midpoint: F = 36.1, p < 0.001), demonstrating strong temporal autocorrelation. We went on to test the hypothesis that SST affected pupping season phenology.

(b) Environmental predictors of phenology

We tested the effects of SST as a predictor of key parameters describing the seal pupping season (table 1). Winter SST was associated with pupping season midpoint: increasing SST was associated with an advance of 7.6 days to the

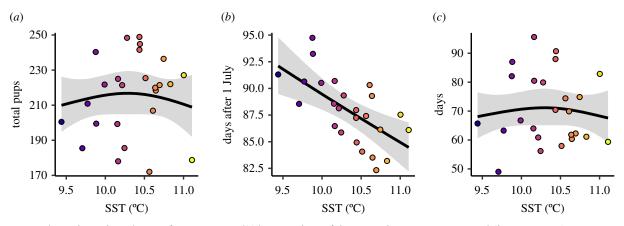


Figure 3. Winter (December—February) sea surface temperature (SST) as a predictor of the grey seal pupping season around Skomer Marine Conservation Zone: the effects of winter SST on (a) total pup count, (b) pupping season midpoint and (c) season length. Solid lines show fitted estimates, with shaded ribbons representing marginal 95% confidence intervals, and points are partial residuals. The fill colour of data points scales with SST. (Online version in colour.)

Table 1. Winter (December—February) sea surface temperature (SST) as a predictor of the grey seal pupping season around Skomer Marine Conservation Zone. (Season parameters are total pups over the season, the midpoint of the season (days since 1 July) and an estimate of 95% of the season length (days).)

	estimates 			SST	
term	min (year)	max (year)	range	F-ratio	<i>p</i> -value
total	166 (2002)	383 (2015)	216	0.001	0.933
midpoint	78 (2004)	99 (1992)	21	2.571	0.018
length	47 (1995)	98 (2011)	51	0.001	0.870

season, explaining 36% of the observed phenological shift (figure 3). However, SST did not explain a statistically significant amount of observed variation in pup numbers or season length.

(c) Individual adult female responses

To understand the phenological patterns that we uncovered, we first explored the prevailing hypothesis; that the mechanism underpinning phenological responses in grey seal pupping operates at the individual adult seal level (through temperature-dependent delayed implantation). Of the adult female seals giving birth to a pup at least once around Skomer, the identities of 150 were recorded (out of a total of 327 adult grey seals identified). Of these, 58 were recorded to have given birth on more than 1 year, and there was no linear trend in the average age of mothers over the monitoring period (F = 3.79, p = 0.084). In those cases, we assessed the role of adult female ageing (estimated by years since first sighting) and SST on date of pupping over successive years. We found no statistical interaction between ageing and SST (Δ AICc = 1.93), suggesting that individual level phenological responses to SST were not dependent on ageing of the mother. Ageing of the mother had a statistically significant effect (figure 4a; F = 9.77, p = 0.002), with adults giving birth to pups earlier as they got older, on average, but SST did not affect when an individual gave birth (figure 4b; F =0.206, p = 0.65). These findings do not support the prevailing hypothesis that sea temperature drives phenological shifts by act directly on individuals' dates of pupping.

(d) Population responses

Next, we tested a novel hypothesis that the phenological mechanism operates at the population level, through modification of the local population age structure. Based on our observations that older females give birth earlier (individual adult dataset), and that warmer SST is associated with earlier pupping seasons (pup population dataset), we predicted that warmer SST would be associated with an increase in the average age of females giving birth (individual adult dataset). As predicted, we found average age of adult females increased with increasing SST (figure 4c; F = 11.8, p = 0.002).

(e) Age-structured population modelling

Our matrix model provided a very good approximation to dynamics, with a finite population growth rate (λ) of 1.07; almost identical to the maximum likelihood estimate of growth rate fitted to empirical population size ($\lambda = 1.07$; electronic supplementary material, figure S3, left panel), and very similar to rates reported elsewhere the UK [73] and North America [74]. As expected, the sensitivity analysis showed that an increase in the transition rates for older breeding adult survival (5-6 and 6+ years) led to an earlier mean pupping date (figure 5a). By contrast, an increase in younger breeding adult survival [4,5] led to a later pupping season, demonstrating the timing of the pupping season is dependent on the relative balance of younger and older breeding adults in the population. However, most importantly, we found that manipulating just these two older stages by increasing

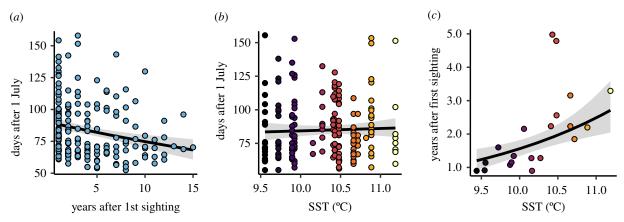


Figure 4. Drivers of grey seal pupping timing at the individual level. The effects of (a) adult female ageing and (b) winter sea surface temperature (SST) on the date of pupping by individual females. (c) The effect of SST on the mean age of adult females in the pupping season at Skomer. Solid lines show fitted estimates, with shaded ribbons representing marginal 95% confidence intervals, and points are partial residuals. The fill colour of data points in (b) and (c) scales with SST. (Online version in colour.)

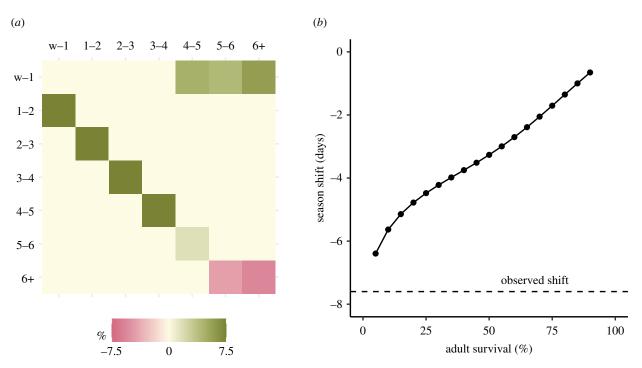


Figure 5. (a) The sensitivity of population-mean pupping date. Pink colours represent a negative sensitivity whereby a positive change in the transition rate leads to an earlier pupping date, whereas green colours indicate a later date. The cream colour represents no change (impossible transitions, in this case). There are two transitions (5–6 and 6+ years) where an increase in transition rate (increased survival) result in a stable age distribution that leads to an earlier mean pupping date. (b) Predicted pupping season shift across a range of possible adult survival values for the 5-6 and 6+ transitions. Adult survival was set at 5% increments between 5% and 90%, then increased to 99.9% and the resulting advance in the pupping season compared to the empirical phenological shift (dashed line). (Online version in colour.)

survival to 99.9% could lead to at most a shift in mean pupping date by just -0.65 days (i.e. 15.7 h earlier).

We further investigated the generality of our MPM by testing a range of breeding adult survival values in the two oldest groups (5-6 and 6+ years), from 5% to our original 90%, and calculating the predicted advance to the pupping season when survival was increased to 0.999. For lower adult survival values, the absolute increase to 99.9% is clearly larger, so a greater advance to the pupping season is predicted. However, even when we set adult breeding survival in the oldest two transitions to just 5%, increasing to 99.9% could only produce a shift of -6.4 days, against our observed advance of 7.6 days (figure 5b).

4. Discussion

While broad-scale, multispecies studies of ecological patterns are critical to identify global challenges resulting from climate change, a recent review has highlighted a lack of mechanistic understanding in this area [12]. This is needed to help us reconcile observed differences in phenological responses to environmental change across, for example latitude, elevation, trophic level, migratory strategy, thermoregulatory mode and generation time. In particular, where research into phenological shifts moves beyond correlative studies of cue and response, these often assume a single mechanism and do not consider or compare alternatives [12]. This is potentially a

critical shortcoming that now presents a barrier to our ability to mitigate against climate change-induced phenological shifts.

We chose a well-studied and regionally important predator species, the grey seal (H. grypus), to test different hypotheses on the underlying mechanism of phenological change. Our key population-level finding was that the peak of the pupping season was more than a week earlier in the warmest years, compared to the coolest. This response was most strongly associated with the previous winter's SST, which we defined as December-February, i.e. SST towards the end of, and shortly after, a given pupping season affects the timing of the following pupping season. This is consistent with the observation that grey seal pupping is earlier in populations that experience warmer SSTs due to latitude [26]. Elsewhere, a negative relationship has also been shown between grey seal birth date and SST, as well as other broad scale climate indicators [32]. The similarity with our population level findings suggests our inferences about the mechanistic basis of this phenological shift may be more widely generalizable.

Our key population-level result is consistent with the mechanism of delayed implantation, set by sea temperatures in the months after mating (which follows autumn pupping in our study population). However, we found no influence of SST in a given year on the timing of pupping for individual mothers. As capital breeders, grey seals accumulate resources for breeding during the majority of the year and then do not forage while suckling their pups [31]. Therefore, the influence of sea conditions earlier in the year is also consistent with resource-limitation affecting competition between individuals. We found that older mothers tended to give birth to pups earlier in the season. This has also been observed in other seal species [58,59], terrestrial mammals [60], as well as many birds [61,62,75].

These observations using population census data provides the basis of an alternative hypothesis based on the age structure of the population: observation (1) warmer years are associated with earlier pupping seasons, and observation (2) older females breed earlier. Therefore, we expected observation (3) the average age of females pupping in warmer years will be greater. We tested this prediction using individual-based, longitudinal data from our focal grey seal population in the SMCZ. Overall, population-level data showed a 1°C increase in observed winter SST equated to around 3–4 days advance in peak pupping date (observation 1). From our individual-level data, we found that a 1-year increase in mean adult female age equated to a 1-day advance in mean pupping date (observation 2). Therefore, we predicted a change of around 6 years to the mean adult female age across the observed temperature range of nearly 2° C. In fact, we observed a difference of just 2 years associated with the observed temperature range (observation 3). However, this discrepancy is within the 95% confidence intervals in parameter estimates so, we believe, represents good support for the age-structured pupping phenology hypothesis.

The SMCZ is a major grey seal pupping site in Wales, so we assume that adult females sighted there are the 'winners' of any competition. Given the time lag with SST, it seems that such competition would be for limiting resources such as prey availability earlier in the year, rather than access to beaches during the pupping season. However, it is known that Welsh seals can travel widely throughout the Irish Sea and as far as southwest England, France and Ireland [30,76–78]. Therefore, a wider study of other pupping sites around the region would be needed to really understand the processes driving dynamics in such a mobile and open population.

Grey seals are particularly well studied in the UK, and similar data to ours have been collected at several Welsh pupping sites, e.g. [79]. It would be possible to infer connectivity between sites using time-lagged spatial cross-correlation of annual pup censuses. However, to fully separate the effects of local demography from movement, direct observation of adult resightings quantifying breeding female site fidelity between years are required. In the Welsh context, the recently published EIRPHOT database of individual adult grey seals, identified from photos, across the Irish Sea would be ideal [30].

The reasons for older females of many species breeding earlier have been explored elsewhere, e.g. [80,81]. This, and associated increases in reproductive success in older females, are hypothesized to result from one or more of: (i) 'selection'—the disappearance of late-breeders over age classes, (ii) 'restraint'—the prioritization of current breeding over future survival in older females, (iii) 'constraint'-improvement in breeding skills with age, and (iv) 'delayed breeding'—the incorporation of early breeders over age classes [82]. Whichever mechanism accounts for an increase in the average age of female grey seals around the SMCZ in warmer years, it is also well documented that older grey seals are typical of long-lived animals, e.g. red deer [83], in having increased success in raising a pup to weaning [31]. Concomitantly, pups born early in the season are more likely to be weaned successfully [84], having potentially important consequences for population resilience and viability with climate change.

We concluded our study by developing an age-structured MPM with which to explore dynamics. This model was motivated by the life history of grey seals and parameterized using grey seal demography data where appropriate [31,85,86]. While our empirical data showed a shift in phenology and age-structure associated with SST, there was no clear, monotonic trend in SST during the study period. Therefore, we question whether SST is acting on the stable age distribution of the population, or through transient effects. If SST affects stable age distribution, then this could affect longterm population growth rate and resilience, since older females are known to have greater weaning success rates [31]. Alternatively, the SMCZ grey seal population is known to mix with neighbouring populations [30], such that SST might change population structure through immigration and emigration. Our MPM clearly showed that biasing the stable age distribution to its limit could not fully account for the observed advance in mean pupping date. This supports the hypothesis that transient changes to the age structure of the population, in response to environmental stochasticity in SST, are currently underpinning observed phenological shifts.

The relationship between age structure, climate change and phenology has been studied across a broad taxonomic range of long-lived animal species, including birds [51,52] and fishes [53], and proposed in mammals [35,87]. The hypothesized underlying mechanism typically focusses on age-dependent differences in local demographic plasticity. For example, in superb fairy wrens (*Malurus cyaneus*), older males are thought to risk moulting in less favourable times of year than younger males [51], in Eurasian blue tits (*Cyanistes caeruleus*) older females are less plastic in their reproductive phenology than younger birds [52], and in walleye pollock (*Gadus chalcogrammus*) climate variation interacts with the age-diversity of the stock to drive mean spawning date [53]. In parallel, climate is known to affect the phenology of long-range movement in

numerous species, which itself may be age-dependent [88-90]. However, our findings indicate that these two research areas, demography and movement ecology, need to be considered together to understand the interplay between population dynamics, climate change and phenology.

In summary, the hypothesis that climate variability acts at the population level, through modification of age structure, provides an alternative to the longstanding, individual-level hypothesis of delayed implantation in large carnivores [35,40]. Moreover, this age structuring mechanism results in population dynamic responses to climate warming that the widely assumed delayed implantation mechanism would not. This may well explain the poor support for a direct link between sea temperatures and pupping phenology in grey seals to date [32,49,50]. More than simply demonstrating a newly recognized phenomenon in a single species, our findings provide motivation and guidance for researchers to consider alternative mechanisms for phenological cueresponse shifts in other long-lived species; in particular, including mechanisms acting at the population level.

Ethics. Spray-marking of seals is conducted under published protocols (Alexander M. 2015 Skomer MCZ and Skomer Island Grey Seal management plan; Poole J. 1996 Grey Seal Monitoring Handbook, Skomer Island. Countryside Council for Wales). Contracted surveyors are issued permits by Natural Resources Wales.

Data accessibility. All data used in this manuscript are available as electronic supplementary material [91].

Authors' contributions. J.C.B.: conceptualization, formal analysis, investigation, methodology, project administration, software, supervision, validation, visualization, writing-original draft, writing-review and editing; O.R.J.: conceptualization, formal analysis, investigation, methodology, software, visualization, writing-original draft, writing-review and editing; L.B.: conceptualization, writingreview and editing; N.F.: data curation, validation, writing-review and editing; R.B.: data curation, validation, writing—review and editing; K.L.: data curation, investigation, methodology, writing—review and editing; T.B.S.: conceptualization, funding acquisition, project administration, resources, writing-review and editing. All authors gave final approval for publication and agreed to be held accountable for the work performed therein.

Competing interests. We declare we have no competing interests.

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References

- 1. Root TL, Price JT, Hall KR, Schneider SH, Rosenzweig C, Pounds JA. 2003 Fingerprints of global warming on wild animals and plants. Nature 421, 57-60. (doi:10.1038/nature01333)
- Menzel A et al. 2006 European phenological response to climate change matches the warming pattern. Glob. Change Biol. 12, 1969-1976. (doi:10. 1111/j.1365-2486.2006.01193.x)
- Møller AP. 2015 Environmental indicators of climate change: phenological aspects. In Environmental indicators (eds RH Armon, O Hänninen), pp. 39-49. Dordrecht: Springer Netherlands.
- Sydeman WJ, Poloczanska E, Reed TE, Thompson SA. 2015 Climate change and marine vertebrates. Science **350**, 772–777. (doi:10.1126/science. aac9874)
- Gallinat AS, Primack RB, Wagner DL. 2015 Autumn, the neglected season in climate change research. Trends Ecol. Evol. 30, 169-176. (doi:10.1016/j.tree. 2015.01.004)
- 6. Brown CJ et al. 2016 Ecological and methodological drivers of species' distribution and phenology responses to climate change. Glob. Change Biol. 22, 1548-1560. (doi:10.1111/gcb.13184)
- 7. Thackeray SJ et al. 2016 Phenological sensitivity to climate across taxa and trophic levels. Nature 535, 241-245. (doi:10.1038/nature18608)
- Mendoza I, Peres CA, Morellato LPC. 2017 Continental-scale patterns and climatic drivers of fruiting phenology: a quantitative Neotropical review. Glob Planet Change 148, 227-241. (doi:10. 1016/j.gloplacha.2016.12.001)
- Piao S et al. 2019 Plant phenology and global climate change: current progresses and challenges.

- Glob. Change Biol. 25, 1922-1940. (doi:10.1111/ acb.14619)
- 10. Visser ME, Both C. 2005 Shifts in phenology due to global climate change: the need for a yardstick. Proc. R. Soc. B 272, 2561–2569. (doi:10.1098/rspb. 2005.3356)
- 11. Miller-Rushing AJ, Høye TT, Inouye DW, Post E. 2010 The effects of phenological mismatches on demography. Phil. Trans. R. Soc. B 365, 3177-3186. (doi:10.1098/rstb.2010.0148)
- 12. Chmura HE, Kharouba HM, Ashander J, Ehlman SM, Rivest EB, Yang LH. 2019 The mechanisms of phenology: the patterns and processes of phenological shifts. Ecol. Monogr. 89, e01337. (doi:10.1002/ecm.1337)
- 13. Miller-Rushing AJ, Weltzin J. 2009 Phenology as a tool to link ecology and sustainable decision making in a dynamic environment. New Phytol. 184, 743-745. (doi:10.1111/j.1469-8137.2009.03083.x)
- 14. Nakazawa T, Doi H. 2012 A perspective on match/ mismatch of phenology in community contexts. *Oikos* **121**, 489–495. (doi:10.1111/j.1600-0706. 2011.20171.x)
- 15. Butt N, Seabrook L, Maron M, Law BS, Dawson TP, Syktus J, Mcalpine CA. 2015 Cascading effects of climate extremes on vertebrate fauna through changes to low-latitude tree flowering and fruiting phenology. Glob. Change Biol. 21, 3267-3277. (doi:10.1111/qcb.12869)
- 16. Renner SS, Zohner CM. 2018 Climate change and phenological mismatch in trophic interactions among plants, insects, and vertebrates. Annu. Rev. Ecol. Evol. Syst. 49, 165-182. (doi:10.1146/annurevecolsys-110617-062535)

- 17. Kudo G, Ida TY. 2013 Early onset of spring increases the phenological mismatch between plants and pollinators. Ecology 94, 2311-2320. (doi:10.1890/ 12-2003.1)
- 18. Conversi A et al. 2015 A holistic view of marine regime shifts. Phil. Trans. R. Soc. B 370, 20130279. (doi:10.1098/rstb.2013.0279)
- 19. Stevenson TJ et al. 2015 Disrupted seasonal biology impacts health, food security and ecosystems. Proc. R. Soc. B 282, 20151453. (doi:10.1098/rspb. 2015.1453)
- 20. Alford RA. 1989 Variation in predator phenology affects predator performance and prey community composition. *Ecology* **70**, 206–219. (doi:10.2307/ 1938427)
- 21. Logan JD, Wolesensky W, Joern A. 2006 Temperature-dependent phenology and predation in arthropod systems. Ecol. Model. 196, 471-482. (doi:10.1016/j.ecolmodel.2006.02.034)
- 22. Yang LH, Rudolf VHW. 2010 Phenology, ontogeny and the effects of climate change on the timing of species interactions. *Ecol. Lett.* **13**, 1–10. (doi:10. 1111/j.1461-0248.2009.01402.x)
- 23. Boyd JM, Campbell RN. 1971 The grey seal (Halichoerus grypus) at North Rona, 1959 to 1968. J. Zool. 164, 469-512. (doi:10.1111/j.1469-7998. 1971.tb01331.x)
- 24. Harwood J, Prime JH. 1978 Some factors affecting the size of British grey seal populations. J. Appl. Ecol. 15, 401-411. (doi:10.2307/2402600)
- 25. Boness DJ, James H. 1979 Reproductive behaviour of the grey seal (Halichoerus grypus) on Sable Island, Nova Scotia. J. Zool. 188, 477-500. (doi:10.1111/j. 1469-7998.1979.tb03430.x)

- 26. Coulson JC. 1981 A study of the factors influencing the timing of breeding in the grey seal Halichoerus grypus. J. Zool. 194, 553-571. (doi:10.1111/j.1469-7998.1981.tb04601.x)
- 27. Hammill MO, Stenson GB, Myers RA, Stobo WT. 1998 Pup production and population trends of the grey seal (Halichoerus grypus) in the Gulf of St. Lawrence. Can. J. Fish. Aquat. Sci. 55, 423-430. (doi:10.1139/f97-218)
- 28. Bowen WD, McMillan J, Mohn R. 2003 Sustained exponential population growth of grey seals at Sable Island, Nova Scotia. ICES J. Mar Sci. 60, 1265-1274. (doi:10.1016/S1054-3139(03)00147-4)
- 29. Russell DJF, Morris CD, Duck CD, Thompson D, Hiby L. 2019 Monitoring long-term changes in UK grey seal pup production. Aquat. Conserv. Mar. Freshw. Ecosyst. **29**, 24–39. (doi:10.1002/agc.3100)
- 30. Langley I et al. 2020 Site use and connectivity of female grey seals (Halichoerus grypus) around Wales. Mar. Biol. 167, 1-15. (doi:10.1007/s00227-020-03697-8)
- 31. Bowen WD, Iverson SJ, Mcmillan JI, Boness DJ. 2006 Reproductive performance in grey seals: age-related improvement and senescence in a capital breeder. J. Anim. Ecol. 75, 1340-1351. (doi:10.1111/j.1365-2656.2006.01157.x)
- 32. Bowen WD, Heyer Cd, Lang SLC, Lidgard D, Iverson SJ. 2020 Exploring causal components of plasticity in grey seal birthdates: effects of intrinsic traits, demography, and climate. Ecol. Evol. **10**, 11 507–11 522. (doi:10.1002/ece3.6787)
- 33. Weimerskirch H, Inchausti P, Guinet C, Barbraud C. 2003 Trends in bird and seal populations as indicators of a system shift in the Southern Ocean. Antarct. Sci. 15, 249-256. (doi:10.1017/ S0954102003001202)
- 34. Learmonth JA, MacLeod CD, Santos MB, Pierce GJ, Crick HQP, Robinson RA. 2006 Potential effects of climate change on marine mammals. Oceanogr. Mar. Biol. Annu. Rev. 44, 431-464.
- 35. Reijnders PJH, Brasseur SMJM, Meesters EHWG. 2010 Earlier pupping in harbour seals, Phoca vitulina. Biol. Lett. 6, 854-857. (doi:10.1098/rsbl. 2010.0468)
- 36. Osinga N, Pen I, Haes Hd, Brakefield PM. 2012 Evidence for a progressively earlier pupping season of the common seal (Phoca vitulina) in the Wadden Sea. J. Mar. Biol. Assoc. UK 92, 1663-1668. (doi:10. 1017/S0025315411000592)
- 37. Cordes LS, Thompson PM. 2013 Variation in breeding phenology provides insights into drivers of long-term population change in harbour seals. Proc. R. Soc. B 280, 20130847. (doi:10.1098/rspb. 2013.0847)
- 38. Gibbens J, Arnould JPY. 2009 Interannual variation in pup production and the timing of breeding in benthic foraging Australian fur seals. Mar. Mammal Sci. 25, 573-587. (doi:10.1111/j.1748-7692.2008. 00270 x)
- 39. Michelle S, Adams G, McCorkell R, Kirkham A, Burns J. 2017 Weddell seal reproductive phenology challenges the notion that all pinnipeds have embryonic diapause. In XIIth SCAR Biology

- Symposium Book of Abstracts, p. 297. See https:// kuleuvencongres.be/scarbiology2017/files/ bookofabstracts.pdf#page=298.
- Jemison LA, Kelly BP. 2001 Pupping phenology and demography of harbor seals (Phoca vitulina richardsi) on Tugidak Island, Alaska. Mar. Mammal *Sci.* **17**, 585–600. (doi:10.1111/j.1748-7692.2001. tb01006.x)
- 41. Tapia-Harris C, Heckel G, Schramm Y, Fernández-Martín EM. 2017 Molting phenology of the Pacific harbor seal (Phoca vitulina richardii) on two islands off the Baja California Peninsula, Mexico. Mar. Mammal Sci. 33, 817-829. (doi:10.1111/mms. 12400)
- 42. Daniel JJ. 1981 Delayed implantation in the northern fur seal (Callorhinus ursinus) and other pinnipeds. J. Reprod. Fertil. Suppl. 29,
- 43. Scheffer VB, York AE, 1997 Timing of implantation in the northern fur seal, Callorhinus ursinus. *J. Mammal.* **78**, 675–683. (doi:10.2307/1382919)
- 44. Laws RM, Baird A, Bryden MM. 2003 Breeding season and embryonic diapause in crabeater seals (Lobodon carcinophagus). Reprod-Camb. 126, 365-370. (doi:10.1530/rep.0.1260365)
- 45. Daniel JC. 1970 Dormant embryos of mammals. BioScience 20, 411-415. (doi:10.2307/1295231)
- Renfree MB. 1978 Embryonic diapause in mammals—a developmental strategy. In *Dormancy* and developmental arrest: experimental analysis in plants and animals (ed. ME Clutter), pp. 1-46. New York, NY: Academic Press Inc.
- 47. Mead RA. 1989 The physiology and evolution of delayed implantation in carnivores. In Carnivore behavior, ecology, and evolution (ed. JL Gittleman), pp. 437-464. Berlin, Germany: Springer.
- 48. Fenelon JC, Renfree MB. 2018 The history of the discovery of embryonic diapause in mammals. Biol. Reprod. 99, 242–251. (doi:10.1093/biolre/ioy112)
- 49. Boyd IL. 1984 The relationship between body condition and the timing of implantation in pregnant grey seals (Halichoerus grypus). J. Zool. 203, 113-123. (doi:10.1111/j.1469-7998.1984. tb06048.x)
- 50. Boyd IL. 1991 Environmental and physiological factors controlling the reproductive cycles of pinnipeds. Can. J. Zool. 69, 1135-1145. (doi:10. 1139/z91-162)
- 51. Pol Mvd, Osmond HL, Cockburn A. 2012 Fluctuations in population composition dampen the impact of phenotypic plasticity on trait dynamics in superb fairy-wrens. J. Anim. Ecol. 81, 411–422. (doi:10.1111/j.1365-2656.2011.01919.x)
- 52. Bonamour S, Chevin L-M, Réale D, Teplitsky C, Charmantier A. 2020 Age-dependent phenological plasticity in a wild bird. J. Anim. Ecol. 89, 2733-2741. (doi:10.1111/1365-2656.13337)
- 53. Rogers LA, Dougherty AB. 2019 Effects of climate and demography on reproductive phenology of a harvested marine fish population. Glob. Change Biol. **25**, 708–720. (doi:10.1111/gcb.14483)
- 54. Büche B, Stubbings E. 2019 Grey seal breeding census Skomer Island 2018. 93. NRW Evidence

- Report Number 325. Aberkenfig, UK: The Wildlife Trust South and West Wales.
- 55. Lock K, Newman P, Burton M, Jones J, Skomer Marine Conservation Zone. 2017 Grey seal survey, Marloes Peninsula 1992-2016. NRW Evidence Report 195. Cardiff, UK: Natural Resources Wales.
- 56. Radford PJ, Summers CF, Young KM. 1978 A statistical procedure for estimating grey seal pup production from a single census. Mammal Rev. **8**, 35–42. (doi:10.1111/j.1365-2907.1978. tb00214.x)
- 57. Austin D, Bowen WD, McMillan Jl. 2004 Intraspecific variation in movement patterns: modeling individual behaviour in a large marine predator. Oikos 105, 15-30. (doi:10.1111/j.0030-1299.1999. 12730.x)
- 58. Reiter J, Panken KJ, Le Boeuf BJ. 1981 Female competition and reproductive success in northern elephant seals. Anim. Behav. 29, 670-687. (doi:10. 1016/S0003-3472(81)80002-4)
- 59. Boyd IL. 1996 Individual variation in the duration of pregnancy and birth date in Antarctic fur seals: the role of environment, age, and sex of fetus. J Mammal. 77, 124-133. (doi:10.2307/
- 60. Plard F, Gaillard J-M, Coulson T, Hewison AJM, Delorme D, Warnant C, Nilsen EB, Bonenfant C. 2014 Long-lived and heavier females give birth earlier in roe deer. Ecography 37, 241-249. (doi:10. 1111/j.1600-0587.2013.00414.x)
- 61. Sydeman WJ, Penniman JF, Penniman TM, Pyle P, Ainley DG. 1991 Breeding performance in the Western Gull: effects of parental age, timing of breeding and year in relation to food availability. J. Anim. Ecol. 60, 135-149. (doi:10.2307/5450)
- 62. Forest Ld, Gaston AJ. 1996 The effect of age on timing of breeding and reproductive success in the thick-billed murre. Ecology 77, 1501-1511. (doi:10. 2307/2265547)
- 63. Vincent C, Meynier L, Ridoux V. 2001 Photoidentification in grey seals: legibility and stability of natural markings. *Mammalian* **65**, 363–372. (doi:10.1515/mamm.2001.65.3.363)
- 64. Pinheiro J, Bates D. 2006 Mixed-effects models in S and S-PLUS. New York, NY: Springer science & business media.
- 65. Huang SH, Meng SXMX, Yang YY. 2009 Assessing the goodness of fit of forest models estimated by nonlinear mixed-model methods. Can. J. For. Res. 39, 2418-2436. (doi:10.1139/X09-140)
- 66. Isojunno S, Matthiopoulos J, Evans PGH. 2012 Harbour porpoise habitat preferences: robust spatiotemporal inferences from opportunistic data. Mar. Ecol. Prog. Ser. 448, 155-170. (doi:10.3354/ meps09415)
- 67. Wood SN. 2006 Low-rank scale-invariant tensor product smooths for generalized additive mixed models. Biometrics 62, 1025-1036. (doi:10.1111/j. 1541-0420.2006.00574.x)
- 68. R Core Team. 2020 R: a language and environment for statistical computing. Vienna, Austria: R Foundation for Statistical Computing. See https:// www.R-project.org/.

- Mazerolle MJ, Mazerolle MMJ. 2017 Package 'AlCcmodavg'. R Package. 281. See https://cran.uib. no/web/packages/AlCcmodavg/AlCcmodavg.pdf.
- 70. Wood SN. 2017 *Generalized additive models: an introduction with R.* New York, NY: CRC Press.
- 71. Pierce D, Pierce MD. 2019 Package 'ncdf4'. See https://www.vps.fmvz.usp.br/CRAN/web/packages/ncdf4/ncdf4.pdf.
- 72. Stott I, Hodgson D, Townley S, Ellner S. 2018
 Package 'popdemo'. See https://cran.r-project.org/
 web/packages/popdemo/popdemo.pdf.
- Summers CF. 1978 Trends in the size of British grey seal populations. *J. Appl. Ecol.* 15, 395–400. (doi:10.2307/2402599)
- 74. Bowen WD, McMillan JI, Blanchard W. 2007 Reduced population growth of gray seals at sable Island: evidence from pup production and age of primiparity. *Mar. Mammal Sci.* **23**, 48–64. (doi:10. 1111/j.1748-7692.2006.00085.x)
- Ezard THG, Becker PH, Coulson T. 2007 Correlations between age, phenotype, and individual contribution to population growth in common terns. *Ecology* 88, 2496–2504. (doi:10.1890/06-2020.1)
- Cronin M, Gerritsen H, Reid D, Jessopp M. 2016 Spatial overlap of grey seals and fisheries in Irish waters, some new insights using telemetry technology and VMS. PLoS ONE 11, e0160564. (doi:10.1371/journal.pone.0160564)
- 77. Vincent C et al. 2017 Grey and harbour seals in France: distribution at sea, connectivity and trends in abundance at Haulout sites. Deep Sea Res Part II

- *Top. Stud. Oceanogr.* **141**, 294–305. (doi:10.1016/j. dsr2.2017.04.004)
- Carter MI et al. 2020 Habitat-based predictions of at-sea distribution for grey and harbour seals in the British Isles: Report to BEIS, OESEA-16-76, OESEA-17-78. St Andrews, UK: Sea Mammal Research Unit.
- Morgan LH, Morris CW, Stringell TB. 2018 Grey seal pupping phenology on Ynys Dewi/Ramsey Island, Pembrokeshire, p. 24. (NRW Evidence Report). Report No.: 156. Cardiff, UK: Natural Resources Wales.
- 80. Curio E. 1983 Why de young birds reproduce less well? *Ibis* **125**, 400–404. (doi:10.1111/j.1474-919X. 1983.tb03130.x)
- 81. Blas J, Hiraldo F. 2010 Proximate and ultimate factors explaining floating behavior in long-lived birds. *Horm. Behav.* **57**, 169–176. (doi:10.1016/j. yhbeh.2009.10.010)
- Goutte A, Antoine É, Weimerskirch H, Chastel O. 2010 Age and the timing of breeding in a long-lived bird: a role for stress hormones? *Funct. Ecol.* 24, 1007–1016. (doi:10.1111/j.1365-2435.2010. 01712.x)
- 83. Guinness FE, Clutton-Brock TH, Albon SD. 1978
 Factors affecting calf mortality in red deer (*Cervus elaphus*). *J. Anim. Ecol.* **47**, 817–832. (doi:10.2307/3673)
- Pomeroy PP, Fedak MA, Rothery P, Anderson S. 1999
 Consequences of maternal size for reproductive expenditure and pupping success of grey seals at North Rona, Scotland. J. Anim. Ecol. 68, 235–253. (doi:10.1046/j.1365-2656.1999.00281.x)

- 85. Lonergan M, Thompson D, Thomas L, Duck C. 2011
 An approximate Bayesian method applied to
 estimating the trajectories of four British grey seal
 (Halichoerus grypus) populations from pup counts.
 J. Mar. Biol. 2011, e597424. (doi:10.1155/2011/597424)
- Thomas L, Russell DJF, Duck CD, Morris CD, Lonergan M, Empacher F, Thompson D, Harwood J. 2019 Modelling the population size and dynamics of the British grey seal. *Aquat. Conserv. Mar. Freshw. Ecosyst.* 29, 6–23. (doi:10.1002/aqc.3134)
- 87. Loe LE *et al.* 2005 Climate predictability and breeding phenology in red deer: timing and synchrony of rutting and calving in Norway and France. *J. Anim. Ecol.* **74**, 579–588. (doi:10.1111/j. 1365-2656.2005.00987.x)
- 88. Rivalan P, Frederiksen M, Lols G, Julliard R. 2007 Contrasting responses of migration strategies in two European thrushes to climate change. *Glob. Change Biol.* **13**, 275–287. (doi:10.1111/j.1365-2486.2006. 01290.x)
- 89. Lehikoinen E, Sparks TH. 2010 Changes in migration. *Eff. Clim. Change Birds* **12**, 89–112.
- Peer AC, Miller TJ. 2014 Climate change, migration phenology, and fisheries management interact with unanticipated consequences. *North Am. J. Fish Manage.* 34, 94–110. (doi:10.1080/02755947.2013. 847877)
- Bull JC, Jones OR, Börger L, Franconi N, Banga R, Lock K, Stringell TB. 2021 Climate causes shifts in grey seal phenology by modifying age structure. Figshare.