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**Environment-sensitive mass changes influence breeding frequency in a capital breeding marine top predator**

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**20 Abstract**

21 1. The trade-off between survival and reproduction in resource-limited iteroparous animals  
22 can result in some individuals missing some breeding opportunities. In practice, even with  
23 the best observation regimes, deciding whether ‘missed’ years represent real pauses in  
24 breeding or failures to detect breeding can be difficult, posing problems for the estimation  
25 of individual reproductive output and overall population fecundity.

26 2. We corrected fecundity estimates by determining if breeding had occurred in skipped  
27 years, using long term capture-recapture observation datasets with parallel longitudinal  
28 mass measurements, based on informative underlying relationships between individuals’  
29 mass, breeding status and environmental drivers in a capital breeding phocid, the grey seal.

30 3. Bayesian modelling considered interacting processes jointly: temporal changes in a  
31 phenotypic covariate (mass); relationship of mass to breeding probability; effects of  
32 maternal breeding state and mark type on resighting. Full reproductive histories were  
33 imputed, with the status of unobserved animals estimated as breeding or non-breeding,  
34 accounting for local environmental variation. Overall fecundity was then derived for Scottish  
35 breeding colonies with contrasting pup production trends.

36 4. Maternal mass affected breeding likelihood. Mothers with low body mass at the end of  
37 breeding were less likely to bear a pup the following year. Successive breeding episodes  
38 incurred a cost in reduced body mass which was more pronounced for North Rona, Outer  
39 Hebrides (NR) mothers. Skipping breeding increased subsequent pupping probability  
40 substantially for low mass females. Poor environmental conditions were associated with  
41 declines in breeding probability at both colonies. Seal mass gain between breeding seasons  
42 was: (i) negatively associated with lagged North Atlantic Oscillation for seals at NR, ; (ii)  
43 positively associated with an index of seal prey (*Ammodytes* spp) abundance at Isle of May,

44 Firth of Forth (IM). Overall fecundity was marginally greater at IM (increasing/stable pup  
45 production) than at NR (decreasing). No effects of mass were detected on maternal survival.  
46 5. Skipping breeding in female grey seals appears to be an individual mass-dependent  
47 constraint moderated by previous reproductive output and local environmental conditions.  
48 Different demographic trends at breeding colonies were consistent with the fecundities  
49 estimated using this method, which is general and adaptable to other situations.

**50 Introduction**

51

52 Resource limitation results in animals making trade-offs between traits such as survival and  
53 reproduction: costly expenditure on breeding can impact future survival or reproduction  
54 (Stearns, 1992). Iteroparous long-lived species may trade off current and future  
55 reproduction when a reproductive episode is sacrificed in favour of survival. True capital  
56 breeders must acquire resources in advance of a breeding attempt to provision themselves  
57 and their offspring throughout the breeding season. Fasting during reproduction means that  
58 maternal resources are finite. Individuals may differ in their ability to acquire resources  
59 according to age or experience (Beauplet, Barbraud, Dabin, Küssener & Guinet, 2006;  
60 Desprez, Pradel, Cam, Monnat & Gimenez, 2011); individual quality (Hamel, Cote, Gaillard  
61 & Festa-Bianchet, 2009) or population density (Hamel, Côté & Festa-Bianchet, 2010).  
62 Moreover, environmental conditions may lead to changing resource availability and  
63 individual life history schedules are more likely to feature missed breeding attempts when  
64 conditions are unfavourable (Cubaynes, Doherty, Schreiber & Gimenez, 2011; Forcada,  
65 Trathan & Murphy, 2008; Parsons 2008; Soldatini, Albores-Barajas, Massa & Gimenez,  
66 2016). Skipped breeding episodes may allow individuals to maintain survival and together  
67 these impact Lifetime Reproductive Output. Population fecundity is affected when sufficient  
68 such events occur.

69

70 The probability of a seal giving birth to a pup is associated with individual body condition,  
71 which results from foraging success (Guinet, Roux, Bonnet & Mison 1998; Stenson, Buren &  
72 Koen-Alonso 2016; Ferguson *et al.* 2017). The mass of a mature female at the start of

73 lactation is a proxy for body condition which can vary between years but also sets limits on  
74 maternal expenditure (the net change in maternal mass from birth to weaning of the pup) in  
75 phocid seals (Arnbom, Fedak & Boyd 1997). Few studies have examined the consequences  
76 of breeding expenditures between seasons for individuals in capital breeding species. Grey  
77 seal (*Halichoerus grypus*) mothers expending substantial resources in one breeding season  
78 were less likely to return to breed in the subsequent year (Pomeroy, Fedak, Rothery &  
79 Anderson 1999). Therefore a mother's mass in one season may influence the probability  
80 that she breeds in the subsequent year, depending on her ability to regain condition  
81 through successful foraging. In other marine predators, biotic and/or abiotic environmental  
82 fluctuations can influence foraging success by changing prey availability, with consequences  
83 for predator condition, reproductive success and survival (Frederiksen, Lebreton, Pradel,  
84 Choquet & Gimenez, 2014; King, Brooks, Morgan & Coulson, 2005; Bost et al., 2015,  
85 McMahon, Harcourt Burton, Daniel & Hindell, 2017).

86

87 Overall, the UK grey seal population has increased in recent decades. Grey seals breed at  
88 approximately 60 colonies in Scotland and the long term decline in pup production at North  
89 Rona (NR) in the Outer Hebrides contrasts with the growth and stabilization seen at the Isle  
90 of May (IM) in the North Sea, reflecting wider regional variation between grey seal breeding  
91 colonies (Duck & Morris, 2016; Smout, King & Pomeroy, 2011a). Currently, an age-  
92 structured population model is fitted to pup production data using Bayesian methods to  
93 estimate grey seal abundance in UK waters (Thomas *et al.*, in press). This model is applied to  
94 wide geographical areas, each of which includes many breeding colonies. As the model's fit  
95 and output are sensitive to prior assumptions concerning vital rates, the accuracy, precision

96 and representativeness of estimates used to inform the priors of the population model are  
97 of fundamental importance, with fecundity a key parameter (Øigård, Frie, Nilssen &  
98 Hammill 2012).

99

100 Determining how often animals breed can be challenging, requiring information additional  
101 to that needed to estimate survival (Desprez, Gimenez, McMahon, Hindell & Harcourt,  
102 2017). Ideally, breeding events can be compiled from direct observations on known,  
103 representative individuals, occurring over the duration of the animals' lifespan, with  
104 individuals equally detectable and breeding state known without error. The latter is  
105 problematic even for long term studies – if an animal is not observed in a given year but is  
106 resighted later, is this a failure to detect a breeding episode or a non-breeding year for that  
107 animal? In practice determining animals' breeding status when they are not observed at  
108 the main study site is difficult and this has prompted development of statistical methods to  
109 account for uncertain state and individual heterogeneity (Rouan, Gaillard, Guédon and  
110 Pradel 2009; King & McCrea 2014; Desprez *et al.* 2017).

111

112 Here we use long-term data from the grey seal breeding colonies on NR and IM including  
113 state specific capture-mark-recapture records and a time-varying covariate, body mass, to  
114 impute the breeding likelihood of female grey seals in skipped breeding years. Simple  
115 estimates of fecundity based on those animals that are observed to attend the colony in a  
116 given year can give an inflated fecundity rate as non-breeding animals may be less likely to  
117 attend (or even if they attend, be re-sighted at) a breeding colony compared to breeding  
118 animals. Therefore, in order to obtain realistic fecundity estimates, it is important to  
119 consider the breeding status of seals that are not observed, potentially because they are

120 absent from the study colony when observations are collected. Although grey seals are  
121 known to show fidelity to their breeding colonies and philopatry (Pomeroy, Anderson, Twiss  
122 & McConnell 1994; Pomeroy, Smout, Moss, Twiss & King 2010) this remains a challenge,  
123 because the numerous/inaccessible alternative breeding colonies are difficult to monitor  
124 adequately through the breeding season (Harrison *et al.* 2006). We use a Bayesian state-  
125 space approach (Royle 2008; King 2012; King & McCrea 2014; Juez, Aldalur, Herrero,  
126 Galarza & Arizaga 2015), assuming that changes in individual mass depend on the breeding  
127 status of animals, and that the probability of subsequent breeding is dependent on body  
128 mass (Pomeroy *et al.*, 1999). Using an underlying process model for the trajectory of  
129 individual mass over time, we estimate the mass and pupping status of unobserved animals,  
130 and hence obtain overall estimates of fecundity for grey seals breeding at the NR and IM  
131 colonies. Within the same integrated modeling framework we test for associations between  
132 mass, vital rates, and environmental drivers (sandeel abundance index for IM, and the NAO  
133 index for NR) and the predicted impacts of these on individual breeding probability and  
134 colony pup production.

135

## 136 **Methods**

137

### 138 *Study colonies and individuals*

139 NR (59.12° N, 5.83° W) Outer Hebrides is 65 km north-west of Cape Wrath, Scotland. The  
140 pupping season on NR spans mid September to late November (Boyd, Lockie & Hewer 1962;  
141 Hiby *et al.* 2013) with peak pupping in early October. Annual pup production declined from  
142 around 2500 in the 1960s to around 500 at present. IM (56.18°N, 2.55°W) lies at the mouth  
143 of the Firth of Forth, Scotland. Until the 1970's few seals bred there, but annual production

144 was over 2000 pups by the 1990s and has fluctuated around this number since then (Duck &  
 145 Morris 2016). The pupping season is from early October to early December, peaking in early  
 146 November. Researchers were present on NR typically between 25 Sep-4 Nov and on IM  
 147 from 25 Oct-6 Dec.

148

149 Results are based on analysis of data from 584 known adult females at NR covering the  
 150 period 1993-2013; on the IM, data were available for 273 adult females from 1987-2014  
 151 (Table 1). At both colonies, individuals were marked with tags or brands, or identified using  
 152 natural pelage patterns (Smout *et al.*, 2011a). Some animals carried combinations of marks,  
 153 sometimes applied at different times, with new animals added to the data set throughout  
 154 the study period (Pomeroy *et al.* 1999; Smout *et al.* 2011a). Most study animals at NR were  
 155 ‘marked’ as breeding adults, very few tagged pups recruited there (Pomeroy *et al.* 2010). At  
 156 IM, up to 25% of known mothers in the early 2000s onwards had retained tags applied  
 157 when they were pups. Further details of study animals, mark-recapture protocols and the  
 158 Cormack-Jolly-Seber (CJS) model for the mark-recapture process including tag loss are  
 159 reported elsewhere (Pomeroy *et al.* 1994; Smout, King & Pomeroy 2011b; McCrea &  
 160 Morgan 2014).

161

Table 1.	<b>NR</b>	<b>IM</b>
<b>Years of data collection</b>	1993-2013	1987-2014
<b>Number of marked animals</b>	584 (394)	273 (1)
<b>Number of marked animals with associated mass data</b>	210	217

162

163 Table 1: Numbers and categories of study animals at each colony. In the second row are counts of animals in  
 164 the mark-recapture data set, with animals entering the study in different years during the study period at each



165 site (numbers in parentheses represent animals identified by pelage-ID only, while the totals include all  
166 animals including those identified with multiple marks for at least some of the years they were observed.) In  
167 the third row are animals with associated mass data.

168

169 Throughout each breeding season, at each colony, researchers surveyed the main breeding  
170 areas daily and the more outlying areas every 3-4d so that seals were identified as soon as  
171 possible after coming ashore. When birth date was not observed directly, it was estimated  
172 using age-related mass and development characteristics (Kovacs & Lavigne 1986).

173 Mother/pup pairs were captured and weighed twice, as close to the start and end of  
174 lactation as possible to allow estimation of maternal postpartum mass ( $M$ ) directly after the  
175 pup is born, and maternal weaning mass ( $W$ ) at the end of lactation, maternal absence  
176 defined weaning date (protocol in Pomeroy *et al.*, 1999). Average normal lactation duration  
177 was 18d (range 14-23d) and we aimed for a minimum of 10d between captures (typically  
178 days 3 and 15 of 18). The study included seals that bred regularly, as well as intermittent  
179 and rarely-sighted breeders. At both colonies, study animals were originally branded or  
180 flipper-tagged as adults and their pups tagged (Smout *et al.* 2011a); additional pups were  
181 tagged at weaning (Pomeroy *et al.*, 2010). "New" animals were added regularly for  
182 weighing. Age ranges of mothers from each colony were similar (5-35 NR, 6-34 IM: ages  
183 were determined from reading incisor tooth sections, or from resights of recruited tagged  
184 pups, Pomeroy *et al.*, 2010). Not all seals captured had a tooth removed for ageing. At NR  
185 from 1998 as many animals as possible were identified by natural markings, many of these  
186 remained observed but not weighed or aged (Hiby *et al.*, 2013).

187

188 *Environmental correlates*

189 UK grey seals are capital breeders: lactating females fast, relying on body reserves accrued  
190 during the preceding foraging period. Food abundance over the year preceding breeding  
191 was expected to play an important role in determining grey seal breeding success (Pomeroy  
192 *et al.*, 1999). To link between breeding, food abundance and environmental conditions, we  
193 selected environmental correlates *a priori* which were relevant to grey seal regional diets,  
194 assuming that both breeding and foraging occurred within the same respective general  
195 areas for each colony (Hammond & Wilson 2016; Russell *et al.* 2013).

196

197 The North Atlantic Oscillation (NAO) annual winter index offers a broad scale measure of  
198 annual meteorological fluctuations, and it has been associated with vital rates of different  
199 species (Thompson & Ollason 2001; King *et al.* 2005; Sandvik, Erikstad, Barrett and Yoccoz  
200 2005). We used 1-year lagged annual winter NAO (i.e. relating to the winter prior to  
201 breeding) to index food (1-group forage fish, Hammond & Wilson 2016) abundance.

202 In the UK, the sandeel *Ammodytes marinus* is found consistently in grey seal diet samples.

203 This high-energy prey appears to be of particular importance for east coast seal populations  
204 (Cury *et al.* 2011; Hammond & Wilson 2016). Indices of sandeel abundance are available in  
205 some areas of the North Sea including areas close to the IM (ICES 2016). However, direct  
206 estimates of sandeel abundance are not consistently available for west coast areas used by  
207 NR seals and sandeels are also less important in west coast seal diets (Hammond, Hall &  
208 Prime 1994; Hammond & Wilson 2016). Sandeels are an important dietary component for  
209 seals foraging around IM therefore sandeel abundance associated with spring/summer  
210 (when sandeels are in the water column and available to fishing) during the year leading up  
211 to grey seal breeding was used as a covariate for the IM colony (ICES 2016; Hammond &  
212 Wilson 2016) .

213

214 *Analytical framework*

215 Adapting the CJS model, we assumed animals were identified by unique marks during each  
216 breeding season (Lebreton, Burnham, Clobert & Anderson. 1992). Individuals were recorded  
217 as 1 or 0 ('seen' or 'not seen'). Pupping status was recorded similarly. Direct mass  
218 measurements were obtained for some of the individuals in the study. We adopted a  
219 Bayesian state-space modeling approach, offering some important advantages e.g. the user  
220 can include informative priors to constrain the parameter search; and sampling from the  
221 posterior distribution of parameters allows for inference about quantities calculated from  
222 model parameters. Adapting the CJS model for this framework required us to separate the  
223 process model (which includes survival and pupping) from the observation model (animals  
224 may or may not have been observed). However, the level of detail that could be included  
225 was limited, due to the nature and quantity of the data available. Because the link between  
226 phenology, breeding and environment was a primary concern, our modelling focused on  
227 this, explicitly including links from environment to individual mass and pupping history, and  
228 then to vital rates. This contrasts with models that assume time-dependent  
229 survival/breeding/recapture probabilities that are common to all individuals, estimating  
230 them separately for each year. In our approach, processes such as foraging success and  
231 mark loss drove the state of the individual (its mass, breeding status and marks present)  
232 through the study period ( King 2014; King, Morgan, Gimenez & Brooks 2009). The  
233 observation process (whether or not the animal was observed) depended on the state of  
234 the animal including its breeding status and identifying marks. After initial exploratory  
235 analyses the initial CMR model in which vital rates depended on mass was further simplified  
236 by excluding mass-dependence in survival rate for both study systems, as there was little

237 evidence for this dependence structure, and survival was estimated separately for each  
238 colony.

239

240 *Process model*

241

242 Maternal expenditure, represented by mass loss during lactation, was estimated using a  
243 general multiplier  $\beta$  acting on maternal postpartum mass (Pomeroy *et al.*, 1999; Wheatley,  
244 Bradshaw, Harcourt & Hindell, 2008; Figure1) . Thus for a female  $j$  pupping in year  $t$  the  
245 expected mass of a female at the start of the breeding season  $M_{j,t}$  was related to her mass  
246 at the end of breeding season

247

$$248 W_{j,t} \sim N(\beta M_{j,t}, \sigma_W^2)$$

249

250 The true mass of the female was assumed to be Normally distributed around the expected  
251 value, reflecting both the individual variation between females, and observation error in  
252 mass measurement. The mass of a female  $j$  at the end of breeding in year  $t$  influenced mass  
253 at the beginning of breeding in year  $t + 1$  subject to an additional colony and year-specific  
254 environmental effect  $\varepsilon_t$  common to all animals, breeders and non-breeders. The effect of  
255 pregnancy on expected mass gain was estimated by the factor  $\delta$  such that for a pregnant  
256 female pupping in year  $t + 1$ , with the true mass of the female assumed to be Normally  
257 distributed:

258

$$259 M_{j,t+1} \sim N(\delta \varepsilon_t W_{j,t}, \sigma_M^2)$$

260

261

262 Thus combining the above modelling components, for a female pupping in year  $t + 1$  the  
263 expected relationship between end-of-season masses in year  $t$  and  $t + 1$  is given by:

264

$$265 \quad E(W_{j,t+1}) = \varepsilon_t \delta \beta W_{j,t}$$

266

267 For a non-pupping female, there was no effect of pregnancy on mass gain and no lactation,  
268 so  $\delta \beta$  were both set to 1. For these non-breeding animals the expected relationship is  
269 described by:

270

$$271 \quad E(W_{j,t+1}) = \varepsilon_t W_{j,t}$$

272

273 The product  $\delta \beta$  could be interpreted as a general estimate of the ratio between the end-  
274 of-season mass for breeding and non-breeding females. Both constants were estimable  
275 because we observe values of both  $W_{j,t}$  and  $M_{j,t}$  in the data set, allowing direct estimation  
276 of  $\beta$  from data on breeding animals. As previously, maternal masses  $W_{j,t}$   
277 were assumed to be Normally distributed with constant variance.

278

279 The year-dependent mass-gain  $\varepsilon_t$  was modelled as a function of the respective  
280 environmental variable (1 year lagged NAO for NR, and sandeel abundance for IM)  
281 represented here by  $x_t$

282

$$283 \quad \varepsilon_t = a + bx_t$$

284

285 where  $a$  and  $b$  were estimated. If the 95% BCI (Bayesian Credible Interval) around the  
286 estimate for the parameter  $b$  did not include zero, this was taken as evidence for an  
287 association between mass gain and the environmental variable.

288

289 Pupping was treated as a Bernoulli process with underlying probability  $f_{j,t+1}$ . This was  
290 associated with maternal weaning mass  $W_{j,t}$  in the previous year, scaled by the year-effect.

291 A logistic relationship was assumed:

292

$$293 \quad f_{j,t+1} = \frac{\exp(a_p + b_p \varepsilon_t W_{j,t})}{1 + \exp(a_p + b_p \varepsilon_t W_{j,t})}$$

294

295 Colony-specific values for  $a_p$  and  $b_p$  were estimated. If the 95% BCI around the estimate for  
296 the  $b_p$  did not include zero, this was taken as evidence for an association between pupping  
297 probability  $f_{j,t+1}$  and  $W_{j,t}$ . The sign of  $b_p$  indicated the type of association.

298

299 Because we could not distinguish between animals that died and any that permanently  
300 emigrated from the study population we estimated ‘apparent survival’, abbreviated to  
301 ‘survival’ hereafter. Preliminary investigations into the effects of maternal mass on survival  
302 did not find evidence for a strong effect of mass on survival so the model structure was  
303 adjusted and survival was estimated as a constant value for each colony (see S1).

304

305 The model included the possibility that some females, referred to here as ‘transients’, were  
306 available to be seen on only one occasion (Pradel, Hines, Lebreton & Nichols. 1997; Hiby *et*

307 *al.* 2013). We estimated the colony-specific probability  $p_{transient}$  that an animal identified  
308 for the first time was in this category.

309

### 310 *Observation model*

311

312 We estimated distinct parameters  $p_{pup}$  representing the re-sighting probability of breeding  
313 females, and  $p_{no\ pup}$  for non-breeding females. Seals were marked with brands, flipper tags,  
314 and pelage markings. Mark-dependent values of re-sighting probability were estimated  
315 during the model-fitting process for NR and IM separately. The probability of tag loss could  
316 also be estimated, because some animals carried multiple mark types e.g. tags and brands.  
317 Brands and pelage-ID were treated as permanent marks (Smout *et al.* 2011a; S1).

318

### 319 *Estimation*

320

321 The Bayesian fitting algorithm estimated values of female mass where gaps occurred in  
322 series (Figure 1). Similarly, the unknown pupping status of unobserved animals was  
323 estimated based on observed masses in the previous and subsequent years.  
324 Unknown mass values and model parameters were estimated using the freely-available  
325 open source software WinBUGS (Lunn, Spiegelhalter Thomas & Best 2009). A model  
326 description, equations and priors are detailed in Supplementary Material (S1); code is  
327 provided in (S2). Convergence was checked based on visual inspection of plots for multiple  
328 chains and BGR convergence statistics (Gelman *et al.* 2013).

329

### 330 *Checking goodness-of-fit*

331 With a complex data set and substantial missing data, it is difficult to directly estimate  
332 goodness of fit or to use information criteria such as DIC to carry out model comparison  
333 (Celeux, Forbes, Robert & Titterington 2006). To address the question of goodness of fit we  
334 have implemented an approach with the same underlying principle as a Bayesian  $p$ -value,  
335 but focused on the estimation of net fecundity - the quantity of particular interest in this  
336 study. In particular our aim is to compare simulated fecundity rates (conditional on the  
337 initial sighting and observed mass if any) with estimated rates drawn from the posterior  
338 distribution of the parameters (i.e. from the MCMC iterations) to identify whether or not  
339 they are comparable. If the estimates are comparable there is no evidence against the  
340 model; alternatively, if they are systematically different this suggests a lack of absolute  
341 model fit with regard to fecundity. This fecundity rate was then used as the associated  
342 “discrepancy function”: we compared the estimated fecundity from the posterior  
343 distribution with the associated simulated fecundity given these particular parameter  
344 values. This process was repeated for 1000 random draws from the posterior. We then  
345 record the proportion of simulated fecundity rates that were higher than the associated  
346 fecundity rate for that posterior sample. As for a formal Bayesian  $p$ -value, if simulated and  
347 fitted rates are similarly distributed, this proportion is expected to be around 0.5, and this  
348 then indicates a satisfactory correspondence between model and data; whereas a  
349 proportion in the “tails” (e.g. lower or upper 5% quantiles) would indicate a potential lack of  
350 model fit (King *et al.*, 2009).

351

352 *Model predictions for pupping probabilities; the effects of varying environmental conditions*

353



354 To explore the implications of the fitted model, we estimated the effect of environmental  
355 drivers on the probability of pupping. ‘Poor environmental conditions’, chosen from the  
356 original covariate data, were defined as the values of sandeel abundance or NAO that  
357 predicted lowest proportional mass gain. ‘Good conditions’ were those values of  
358 environmental covariates for which predicted proportional mass gain was highest. We  
359 estimated the ‘skipping point maternal weaning mass’  $W_{50}$  as the mass of the mother when  
360 her pup weaned at which pupping probability in the next year would be 50%, if conditions  
361 during the intervening year were poor. This was calculated using the fitted model for each  
362 colony. Similarly the probability of pupping for a female with mass at weaning  $W_{50}$  after a  
363 year of good conditions was also calculated. Therefore, in effect we tested a theoretical  
364 ‘skipping point mother’ against the worst and best conditions observed in the data.

365

366 The consequences of the variable environment, were explored further by estimating  
367 pupping probabilities for females starting at  $W_{50}$  after two years of good conditions, and  
368 after two years of poor conditions. These values were calculated either assuming that  
369 pupping took place in the breeding season after the first year, or that it did not.

370

### 371 *Model predictions for local population trends*

372 To explore the consequences of our model estimates for local populations, we used a simple  
373 Leslie matrix population simulation for females breeding at IM and NR, assuming no density-  
374 dependent effects were acting and accounting for transients. In this model all animals  
375 became fertile aged 6, adult and sub-adult survival rates were set at the mean estimated  
376 values for adults for each colony, and fecundity was set to the mean colony average  
377 estimate. Female first-year survival was set at 0.6 for IM (Hall & McConnell 2007) and for

378 NR this value was set at  $0.6s_{NR}/s_{IM}$  where  $s_{NR}$  and  $s_{IM}$  were the estimated adult survival  
 379 rates at NR and IM.

380

## 381 Results

382

383 After a 2000 iteration burn-in period, the MCMC for NR and IM appeared to converge with  
 384 stationary posterior parameter estimates achieved after 100,000 iterations. Breeding  
 385 parameter estimates are summarized in Table 2, for full parameter table see Supplementary  
 386 Material (S1). The estimated values from our Goodness of Fit tests were 0.307 for NR and  
 387 0.284 for IM which were both satisfactory, indicating acceptable model fit for fecundity rate  
 388 estimation.

### 389 Parameter estimates

Table 2.	Meaning	NR value	IM value
$s$	Overall survival rate (average over all years)	0.89 (0.87, 0.90)	0.94 (0.93, 0.95)
$f$	General estimate of fecundity for all animals at the colony, including years they are not observed at the colony	0.79 (0.77, 0.81)	0.82 (0.79, 0.84)
$\beta$	Ratio of W (maternal mass at weaning) to M (maternal post partum mass)	0.65 (0.64, 0.66)	0.65 (0.65, 0.66)
$\delta$	Maternal mass gain (preg)	1.34 (1.32, 1.36)	1.40 (1.38, 1.42)
$p_{pup}$	Probability that a female marked with a brand (i.e. highly visible) and pupping is present and will be seen at the colony	0.89 (0.85, 0.93)	0.83 (0.80, 0.86)
$p_{no\ pup}$	Probability that an animal marked with a brand which is not pupping will be seen at the colony	0.08 (0.06, 0.11)	0.05 (0.03, 0.08)
$P_{transient}$	Probability that a female recorded in the data set for the first time is a 'transient'	0.21 (0.16, 0.27)	0.04 (0.01, 0.09)
$P_{tagloss}$	Annual probability of tag loss	0.07 (0.05, 0.09)	0.02 (0.01, 0.03)

390

391 Table 2: Mean parameter estimates for the process and observation models (95% Bayesian credible  
 392 intervals are shown in brackets).

393

394 Mean survival rate at NR was lower than at IM, consistent with previous findings (Smout *et*  
395 *al.* 2011a). There were more mothers estimated as ‘transient’ at NR than at IM (Table 2).

396

397 For both colonies there was evidence for a positive relationship between pupping  
398 probability in year  $t + 1$  and  $W_t$  (maternal mass at the end of breeding in year  $t$ ) with  
399 appreciable effects on pupping probability over the range of mass values observed in the  
400 data set (Figure 2). The IM pupping probability-mass curve shows a steeper relationship  
401 than that for NR: at IM, pupping probability doubles from 0.4 to 0.8 over a range of  
402 approximately 13kg while at NR the same change occurs over a range of approximately  
403 23kg.

404

405 Average net fecundity calculated for NR females is less than that for IM females, but  
406 credible intervals overlap for the two sites. For comparison, a simple calculation based on  
407 the observational data, assuming that all unobserved known animal-years were non-  
408 breeding years, gave net fecundity rates of 0.679 at NR and 0.750 at IM, highlighting the  
409 importance of accounting for unobserved individuals.

410

411 The estimated probability of re-sighting a non-breeding female was low at both colonies,  
412 consistent with the observation that very few non-breeding adult females are seen at these  
413 colonies (Table 2).

414

415 The postpartum masses of mothers at NR and IM encompassed a similar range, with several  
416 at each colony exceeding 250kg. The estimated value of  $\beta$  (the ratio of maternal mass at the

417 end of lactation to mass at start of lactation) was very similar between IM and NR: on  
418 average, a mother expended 35% of her postpartum mass on raising a pup (Table 2).  
419 Average proportional mass gain  $\delta$  for pregnant mothers between end and start of  
420 successive breeding seasons was lower for those at NR than for those at IM (Table 2).

421

#### 422 *Environmental variation and breeding probability*

423

424 There was no evidence for an association between mass gain and NAO at IM, so this was  
425 excluded from the final fitted model. The relationship between mass gain and sandeel  
426 abundance at IM was positive (Figure 3). For NR, the underlying relationship between mass  
427 gain and NAO index was negative (Figure 4).

428

429 Year specific proportional mass gains  $\varepsilon_t$  were generally larger at NR than at IM (Figures 3,4;  
430 right panels). Synchronicity of environmental effects at the two colonies was limited; better  
431 than average mass gains occurred at both colonies in 1995 and 2010.

432

433 The predicted effects of environment and breeding on subsequent pupping probability are  
434 important for ‘skipping point’  $W_{50}$  animals at both colonies ( $p_{t+1}$  in Table 3). Not breeding  
435 has an appreciable effect for skipping point animals, where pupping probability  $p_{t+2}$  can  
436 either decrease if pupping takes place in year  $t+1$ , or increase if breeding is skipped (Table 3,  
437 columns 4 and 5). Mothers at the skipping point were heavier at NR (93.5kg) than at IM  
438 (88.5kg).

439

440

Table 3.	Environmental conditions	$p_{t+1}$	$p_{t+2}$ (pupped $t+1$ )	$p_{t+2}$ (skipped $t+1$ )
NR	Poor - High NAO	0.500	0.423	0.696
	Good - Low NAO	0.633	0.675	0.884
IM	Poor - Low sandeels	0.500	0.426	0.709
	Good - High sandeels	0.620	0.647	0.868

441

442 Table 3: The predicted impacts of environmental conditions. ‘Skipping point’ mothers of mass  $W_{50}$  in year  $t$   
 443 have a 50% probability of pupping in year  $t+1$  after a ‘poor’ year of environmental conditions. If  
 444 environmental conditions are better (low NAO at NR, high sandeels at IM) then pupping probability in year  $t+1$   
 445 is improved ( $p_{t+1}$  column 3). Predicted pupping probabilities for year  $t+2$  are shown after 2 years of consistent  
 446 environmental conditions (2 good years or 2 bad years), in columns 4 and 5. Column 4 gives values for females  
 447 if they bred in year  $t+1$ ; Column 5 gives values for females that ‘skipped’ breeding in year  $t+1$ .

448

449 *Population trajectory for each colony*

450

451 The predicted time series of number of females breeding derived from the Leslie model  
 452 declined at NR and increased at IM (dashed and solid lines respectively, Figure 5). Simple  
 453 visual comparison suggests that there is a good correspondence with trends in pup  
 454 production estimates derived from synoptic counts of pups from aerial survey (NR, IM solid  
 455 and open circles respectively, Duck & Morris 2016).

456

## 457 Discussion

458

459 Intermittent breeding in iteroparous animals acts on LRO and individual fitness, and affects  
 460 demographic rates at the population level. Capital breeding species in fluctuating  
 461 environments are likely to be particularly prone to skipping breeding episodes, when  
 462 survival takes precedence over reproduction (Stearns, 1992). Skipped breeding episodes

463 among experienced breeders are known in capital breeding phocid seals, but mechanistic  
464 explanations of the process have often relied anecdotally on Boyd's (2000) finding that body  
465 mass accounted for more variation in pregnancy rates in capital breeding than income  
466 breeding species (e.g. Chambert, Rotella, Higgs & Garrott, 2013; Desprez et al., 2017).  
467 Desprez et al. (2017) used resighting records at Macquarie Island to investigate intermittent  
468 breeding of southern elephant seals *Mirounga leonina*. Using a multi-event model  
469 incorporating uncertain reproductive status and categorizing adult females into breeding  
470 heterogeneity classes, they found that there was a survival cost to breeding for females in  
471 the infrequently breeding class. This latter class they suggested comprised females of lower  
472 quality in some phenotypic property positively related to fitness, and while variability in  
473 individuals' ability to forage and gain resources were hypothesized to be likely explanatory  
474 factors, no phenotypic or other measures of animals were provided to support this (Desprez  
475 et al., 2017). Similarly, Chambert, Rotella & Garrott, (2015) proposed that female Weddell  
476 seals (*Leptonychotes weddellii*) that skipped breeding in Erebus Bay, Antarctica were in  
477 poorer condition than regular breeders. Here, we provide the empirical evidence for an  
478 effect of maternal mass on pupping probability in a capital breeding phocid, the grey seal,  
479 and importantly, of the increased pupping probability that low body mass females accrue by  
480 skipping a reproductive episode.

481

482 Environmental fluctuations are seen to impact long-lived marine species' fecundity rather  
483 than survival (Reed, Harris & Wanless, 2015; Stenson et al., 2016). Our study showed an  
484 impact of female mass on the probability of pupping but did not detect an effect on survival.  
485 While a female in sufficiently poor condition might be expected to have reduced survival  
486 probability, this may be difficult to observe. If low-mass animals miss breeding, are not

487 observed at the colony and are never seen again we may lack the data to define the shape  
488 of the descending arm of the logistic curve for the survival-mass relationship. If breeding is  
489 costly, animals in poor condition (low body mass) may forego breeding and improve their  
490 own chances of survival (Ronget, Garratt, Lemaître & Gaillard 2017). We found that impacts  
491 of breeding on subsequent pupping probability for grey seals are substantial for seals whose  
492 mass at end of breeding is low (Figure 2). At both colonies our model predicts that in poor  
493 environmental conditions, females that ‘skip’ breeding are more likely to pup in subsequent  
494 years (Table 3). Similar skipping breeding sequences were linked with body condition and  
495 environmental conditions in Weddell seals (Proffitt, Garrott, Rotella & Wheatley 2007;  
496 Chambert et al., 2015).

497

498 Mass-adjusted fecundity rates suggest that around two-thirds of “missing” years are indeed  
499 non-breeding years (Table 2). Movement away from the natal colony or permanent  
500 emigration by adults to different colonies might occur, and cannot be distinguished here  
501 from permanent loss of fertility, or death (Harrison et al., 2006). Nevertheless, our analyses  
502 suggest breeding occurred in around one third of “missed” breeding years. Some pupping  
503 events of known females at their “normal” colony could have gone unobserved, but high  
504 resight rates (Table 2) suggest these should be few. The remaining ‘missed’ breeding events  
505 must have occurred outwith the prevalent breeding site fidelity pattern (Pomeroy et al.  
506 1994). Genetic differences between NR and IM seals are sufficient to suggest very low  
507 effective breeding interchange between distant colonies (Allen, Amos, Pomeroy & Twiss,  
508 1995), but any breeding colony infidelity would help explain unassigned paternities  
509 (Worthington Wilmer, Allen, Pomeroy, Twiss & Amos 2003).

510

511 Some female grey seals observed for the first time became subsequently unobservable,  
512 especially at NR (Table 2). Although termed “transients”, different phenomena can produce  
513 such records. Most convincing is the difficulty in making matches between patterns  
514 extracted from different photographs of the same grey seal identified by pelage alone (Hiby  
515 *et al.* 2013), as many more females were identified by this method at NR (Table 1).  
516 Alternative explanations include: some subset of adult females have lower survival e.g.  
517 primiparous females; higher tag loss rates directly after application; different tag loss rates  
518 between seals at different colonies. Seals may visit a colony once only, then breed  
519 elsewhere subsequently - features of a declining colony such as NR may indicate to first time  
520 visitors that it is undesirable, resulting in demographic state-dependent colony fidelity.  
521  
522 Intermittent breeding has been described as a tactic employed by poorer quality mothers to  
523 optimize LRO (Desprez *et al.*, 2017). True capital breeders rely on accumulated reserves,  
524 expending a large proportion of postpartum body mass (grey seals 0.35, Table 2; southern  
525 elephant seals 0.35, Arnborn *et al.*, 1997) to sustain a reproductive episode. Average  
526 proportional expenditure was the same at NR and IM, despite body mass differences and  
527 different vital rates. Pregnant NR seals did not regain as much of their previous MPPM as  
528 equivalent mothers at IM ( $\delta = 1.34, 1.40$  respectively, Table 2), suggesting that successive  
529 pregnancies incur a cost in reduced subsequent MPPM and that NR mothers experienced  
530 this to greater extent, possibly as a result of poorer overall resource availability. Further  
531 reductions in mass of small mothers would be unsustainable and in such cases skipping  
532 breeding seems likely. While a mother experiences strong selective pressure on raising a  
533 pup within resource limits, the future costs of which vary according to initial maternal body  
534 mass and reserves, individual and episodic variations in reproductive expenditure occur



535 (Arnbom et al., 1997; Pomeroy et al., 1999). However, even for capital breeders, body  
536 mass alone may not be a perfect metric of body reserves, because it masks the availability  
537 of all the nutrients required for a successful breeding episode. Often only lipid reserves are  
538 used to estimate “condition” in capital breeders as they provide most of the energetic  
539 requirements of mother and offspring, but protein and other nutrients must also be  
540 available and in the extreme may be more limiting (eg Arnbom et al., 1997; Boyd, 2000;  
541 Mellish, Iverson & Bowen, 1999; Hanson, Smout, Moss & Pomeroy, in press). Our modelling  
542 framework could be adapted to include more complex measures of body condition in future  
543 studies, including combining different measures, if such data are available.

544

545 Assessing the effects of different demographic rates on populations requires that rates are  
546 estimated appropriately: for fecundity, non-breeders and non-breeding episodes must be  
547 taken into account (Lee, Reid & Beissinger 2017). Here, we found that relatively small  
548 differences in colony fecundity rate are associated with very different colony pup  
549 production trajectories. Canadian and Norwegian studies of grey seals have reported  
550 fecundity estimates similar to our mass-adjusted overall female fecundity rates (Bowen,  
551 Iverson, McMillan & Boness 2006; Hammill & Gosselin 1995). Nevertheless, fecundity and  
552 adult female survival were both lower for NR seals than those at the growing IM colony,  
553 which had similar parameters to those reported from the expanding colony on Sable Island,  
554 Canada (den Heyer & Bowen, 2017). Simple Leslie matrix population simulations showed  
555 rates of local population growth and decline that are consistent with observed trends in pup  
556 production at both colonies for the early years of the study: a decline in pup production at  
557 NR, compared with a positive trend at IM (Duck & Morris 2016). The breeding decline at  
558 NR is present at other Hebridean grey seal colonies (Duck & Morris 2016) and poor

559 recruitment of seals tagged as pups to NR is a likely additional factor in explaining these  
560 trends (Pomeroy *et al.*, 2010). Long term declines in measures of grey seal “condition” at  
561 NR compared to IM suggests that colony-level effects are reflected in individual phenotypic  
562 covariates and these are a local response to local conditions (Hanson *et al.*, in press) which  
563 likely explains lower fecundity at NR (Boyd, 2000). Longitudinal seabird surveys on NR have  
564 shown decadal declines in 9 of 15 species that use the island for breeding, in common with  
565 trends reported in such species elsewhere in the North Atlantic (Murray & Wilson 2013).  
566 That many different marine apex predators show contemporaneous regional declines is  
567 indicative of the generality of ecosystem change to their detriment in this region.

568

569 Grey seals are generalist predators and their diet varies in response to changing prey  
570 abundance (Smout, Rindorf, Hammond, Harwood & Matthiopoulos 2013). As capital  
571 breeders capable of wide-ranging foraging, they are buffered against small scale, short term  
572 changes in prey availability. The association between NAO (lagged one year) and grey seal  
573 mass change at NR, is explicable if Atlantic atmospheric conditions correlate with the  
574 abundance of one or more important prey species with direct consequences for the  
575 condition of seals foraging and breeding on the UK’s Atlantic fringe. There was no evidence  
576 of a similar effect of NAO at IM, consistent with the limited influence of NAO on the North  
577 Sea ecosystem where most IM seals forage, compared to west coast waters that are  
578 connected directly to the Atlantic (Harris, Edwards & Olhede 2014). The appreciable effect  
579 of local sandeel abundance reflects the importance of sandeels in seal diets in this area and  
580 the wider North Sea where these seals are likely to forage (Russell *et al.*, 2013; Hammond &  
581 Wilson 2016).

582

583 The Bayesian modelling approach implemented here using freely available software allowed  
584 us to fit a model of the relationship between annual mass gain and environmental drivers,  
585 including ‘nuisance’ processes (tag loss) that might otherwise obscure important effects and  
586 a realistic observation model with missing data and unknown states. Our analyses  
587 highlighted the consequential nature of breeding events: longitudinal maternal mass  
588 trajectories suggested underlying breeding histories, and while skipping breeding was  
589 associated with low body mass, females that skipped had a higher probability of breeding  
590 subsequently.

591

## 592 **Ethics**

593 Work involving animals in this study was licensed under UK Home Office project 60/4009 or  
594 preceding versions and conformed to the UK Animals (Scientific Procedures) Act, 1986.

595 Research was approved by the University of St Andrews Animal Welfare and Ethics  
596 Committee.

## 597 **Data accessibility**

598 Data are lodged in PURE, [http://dx.doi.org/10.17630/8b1f33d5-50bf-4631-9e65-](http://dx.doi.org/10.17630/8b1f33d5-50bf-4631-9e65-fca33d26902f)  
599 [fca33d26902f](http://dx.doi.org/10.17630/8b1f33d5-50bf-4631-9e65-fca33d26902f).

## 600 **Authors' contributions**

601 PP designed and managed the long term research and carried out fieldwork; PP and RK  
602 conceived of and designed the study; SCS analyzed the data; SCS and PP led writing of the  
603 paper and share first authorship. All authors gave final approval for publication.

**604 Competing interests**

605 The authors have no competing interests.

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