



## 25 **Abstract**

26 Weaning marks a critical life-history transition in mammals, shaped by maternal investment  
27 strategies, offspring condition, and environmental constraints. In Galápagos sea lions (*Zalophus*  
28 *wollebaeki*), weaning age is highly variable, with some individuals continuing to suckle into  
29 adulthood (supersucklers). Using 20 years of mark-recapture data from 1890 individuals, we  
30 applied multi-state capture-recapture models to investigate the probability of weaning age, and  
31 logistic regression models to explore the drivers of supersuckling. We examined maternal effects  
32 (age, birth interval, and pup condition at birth), offspring traits (sex, early growth), and  
33 environmental variability (Oceanic Niño Index [ONI]). Weaning probability peaked at four years  
34 of age, with 95% of individuals weaned by this time. Longer birth intervals were associated with  
35 later weaning. Offspring sex had no influence on weaning probability. However, ONI affected  
36 juvenile survival, indirectly shaping weaning outcomes. No intrinsic predictors of supersuckling  
37 were identified, although low resource availability during an individual's second or third year of  
38 life increased the likelihood of becoming a supersuckler. While weaning is typically driven by  
39 escalating mother-offspring conflict, the persistence of supersuckling challenges this paradigm.  
40 We propose that the regular occurrence of supersuckling in this species may therefore be socially  
41 driven, reflecting the maintenance of long-term mother-offspring relationships that contribute to a  
42 complex adult social structure. This study highlights the interplay between environmental  
43 unpredictability and social behaviours in shaping life-history strategies.

44

## 45 **1. Introduction**

46 Weaning in mammals marks the end of maternal nutritional investment and is a fundamental life-  
47 history transition for offspring. The timing of weaning is shaped by a complex interplay of  
48 maternal investment strategies, offspring condition, and environmental constraints (Lee 1996;  
49 Schulz & Bowen 2005; Avery & Zinn 2023). As an energetically costly form of parental care,  
50 lactation must be balanced against the mother's future reproductive success, leading to an inherent  
51 mother-offspring conflict over the duration of care (Trivers 1974). Offspring who benefit from  
52 protracted maternal investment in the form of nutrition may have increased chances of survival  
53 (Gerber et al. 2021; Dittus & Baker 2023; Vitikainen et al. 2023). However, gluttonous offspring,  
54 once weaned, may be unable to maintain the same nutritional input and subsequently perish (Reiter  
55 et al 1978). Mothers, on the other hand, may incur an immediate cost to body condition and

56 reproductive fitness but be rewarded for their extended investment through delayed generational  
57 success (Péron et al. 2019; Zippel et al. 2020; Bleu et al. 2021; Chapman et al. 2021; Nielsen et al  
58 2023).

59 Amongst pinnipeds the duration of lactation is particularly variable due to their terrestrial or floe  
60 ice breeding and aquatic foraging (Pomeroy 2011; Avery & Zinn 2023). Phocids generally exhibit  
61 abrupt weaning after an intensive lactation period, whereas otariids (eared seals) experience  
62 prolonged lactation and a gradual transition to nutritional independence that can last up to 36  
63 months in some species (Sepúlveda & Harcourt 2021). This lengthily dependence increases the  
64 potential for variability in weaning age and may intensify mother-offspring conflict (Trillmich &  
65 Wolf 2008; Sepúlveda & Harcourt 2021; Dittus & Baker 2023). Maternal investment may be  
66 determined by pup traits such as sex, mass and demand (Goldsworthy 2006; McDonald et al. 2012;  
67 Kraus et al. 2013). However, lactation duration, and therefore weaning age, is frequently dictated  
68 by maternal factors such as parity, maternal resources and reproductive experience (Schulz &  
69 Bowen 2004; Trillmich & Wolf 2008; Costa & Maresh 2022). Environmental factors such as  
70 predation (Ferguson 2006) and climatic variability (Burkanov et al. 2011; Costa & Valenzuela-  
71 Toro 2021; Schwarz et al. 2022) further influence weaning by affecting maternal foraging success  
72 and offspring condition. During periods of resource scarcity, females may prioritize their own  
73 survival and future reproductive opportunities, leading to earlier weaning or decreased investment  
74 in current offspring (Soto et al. 2004). Conversely, in times of higher resource availability,  
75 prolonged lactation may be more feasible, allowing offspring to remain nutritionally dependent  
76 for extended periods.

77 Galápagos sea lions (GSL) *Zalophus wollebaeki* are an endangered otariid endemic to the  
78 Galápagos archipelago. As an equatorial species GSL are exposed to extreme stochastic  
79 environmental fluctuations in the form of El Niño events which directly impact resource  
80 availability for foraging mothers (Soto et al. 2004; Schwarz et al. 2022). The reproductive period  
81 is protracted starting in September and ending in January (Pörschmann et al. 2010) although  
82 females show high asynchronicity and low seasonality in their breeding cycle (Villegas-Amtmann  
83 et al. 2009). There are high amounts of individual variation in age at primiparity (4-9 years) and  
84 the time until the birth of the next sibling (birth interval). However, females produce on average,  
85 a single pup every two years from the age of six (Kalberer et al. 2018). The expected lifespan for

86 GSL is 21 years (Krüger et al. 2021) and the oldest recorded age at parturition is 18 years (*personal*  
87 *communication*). Adoption and alloparenting are extremely rare in this species (Childs et al. 2025).  
88 Reproductive overlap is not uncommon and, whilst some females have been observed supporting  
89 multiple dependent offspring, more often than not, the situation results in sibling competition and  
90 the rejection or death of one offspring (Trillmich & Wolf 2008). Recent research has shown that  
91 GSL exhibit a prolonged and flexible lactation period lasting between one and three years (Childs  
92 et al. 2025), with some individuals, ‘supersucklers’ continuing to suckle into adulthood. However,  
93 robust statistical analysis of age at weaning has yet to be conducted.

94 Accurately determining weaning age is crucial for understanding both the physiological and  
95 ecological determinants of maternal investment strategies in otariids (Costa & Valenzuela-Toro  
96 2021; McHuron et al. 2023). However, direct observations of weaning are often logistically  
97 challenging, particularly in long-lived, wide-ranging marine species (Trillmich et al. 2016;  
98 Osiecka et al. 2020; Hastings et al. 2021). Multi-state hidden Markov models (HMMs) provide a  
99 powerful tool for inferring weaning transitions from mark-recapture data, allowing estimates of  
100 state probabilities and the role life-history traits and environmental fluctuation may play in  
101 variability (Hastings et al. 2021).

102 Using multi-state capture-recapture models (MSCRM) framed as HMMs, we investigate weaning  
103 age in GSL in relation to life-history traits and environmental conditions. We predicted that low  
104 resource availability, represented by the Oceanic Niño Index (ONI) anomaly, would delay  
105 weaning. Moreover, we hypothesised that male offspring would wean later due to male-dominant  
106 size dimorphism and that lower maternal investment and inexperience would prolong the lactation  
107 period. Based on the MSCRM results we further explored the possible drivers of supersuckling  
108 using logistic regression models (LRM). We theorise that supersucklers exhibit slower growth than  
109 their non-supersuckling conspecifics and that poor environmental conditions during the year of  
110 expected weaning may lead to delayed weaning. A better understanding of weaning patterns and  
111 the mother-offspring relationship in this species is particularly relevant in the context of increasing  
112 environmental variability, as shifts in resource availability may influence maternal investment  
113 strategies.

114

115 **2. Methods**

116

117 **2.1. Data collection & Study Site**

118 Caamaño islet (0°45'S, 90°16'W) is located off the southern coast of Santa Cruz, Galápagos, and  
119 is home to a breeding colony of GSL that has been intensively studied since 2003 (Wolf &  
120 Trillmich, 2007). Data for this study were collected between February 2003 and April 2024 during  
121 two annual field seasons (October–December and February–April) and grouped by reproductive  
122 season (August–July), referred to by the calendar year in which each season began. All individuals  
123 that are born on, or regularly use the islet are given a pit-tag, and marked using Allflex tags,  
124 resulting in a database of extensive individual life history (for details on capture methodology see  
125 Wolf & Trillmich 2007; Meise et al. 2013). Identification rounds were conducted daily (2-5 times),  
126 during which the ID, location, and suckling events of all individuals present on the islet were  
127 recorded (for details on identification rounds see Trillmich et al. 2016; and for suckling events  
128 Childs et al. 2025). Missing tags were recorded whenever tag-loss was observed and used as means  
129 of tracking an individual's absence from the database. Considerable effort was made to mark  
130 unidentified mothers and replace missing tags to maintain life history records.

131

132 Mother-offspring pairs were identified through direct observation of birth, perinatal bonding, or  
133 repeat suckling events within a month (Childs et al. 2025). Each individuals' date of birth (DoB)  
134 was categorised: where 'exact' is accurate to the day, 'weak' is  $\pm 2$  weeks, and 'unknown' is  
135 greater than 2 weeks uncertainty. Field-based age estimates for juveniles have proven to be highly  
136 accurate (Jeglinski et al. 2010). We therefore recategorised all individuals originally labelled as  
137 'unknown' DOB and first marked at, or under, 6 months of age as 'roughly known' so that they  
138 could be reassigned to their appropriate reproductive year of birth. All individuals born before  
139 2003, when data collection first began, were categorised as DOB 'unknown'. Individuals with  
140 unknown DoB were not included in any statistical modelling or testing involving age.

141

142 Sex verification and morphological measurements were taken every time an individual was  
143 captured. Individuals were weighed with an electric scale, and length was measured from the tip  
144 of the snout to the tip of the tail (see Mueller et al. 2011). All individuals were captured 1. for  
145 marking, 2. to observe pup growth during their first 2 months, 3. annually to measure juvenile  
146 morphological differences and cohort response to environmental variations (Wolf & Trillmich

147 2007). To avoid disturbance to the mother-offspring relationship, offspring are only captured when  
148 their mother is absent. Offspring body condition was calculated using scaled mass index (SMI)  
149 (Peig & Green 2009) for both birth (10-day-old or less) and early life (90-day-old mean) as  
150 indicators of offspring health and maternal investment (Mueller et al. 2011). Which we take as  
151 representative of maternal investment due to the offspring's nutritional dependence on milk at this  
152 life stage (Jeglinski et al. 2012; Schwarz et al. 2022). For investigating growth differences between  
153 supersucklers and non-supersucklers, SMI was calculated for each time point (age) where both  
154 length and weight measurements were available.

155

156 Birth interval, measured as the number of years until the birth of the next sibling, was used to gain  
157 insight into the effect the presence of a younger sibling may have on weaning age as sibling conflict  
158 is not uncommon in this species and females are known to reject older offspring upon the birth of  
159 a new pup (Trillmich & Wolf 2008). Due to the possibility that longer than average intervals (2  
160 years, Kalberer et al. 2018) might be due to tag-loss, unrecorded miscarriages or stillbirths, females  
161 not observed on the islet for more than 2 years were excluded from the sample. We further cross-  
162 referenced all individuals with known multiple-year tag-loss to check for mothers who may have  
163 given birth, but no offspring was assigned due to an inability to positively identify the pair.

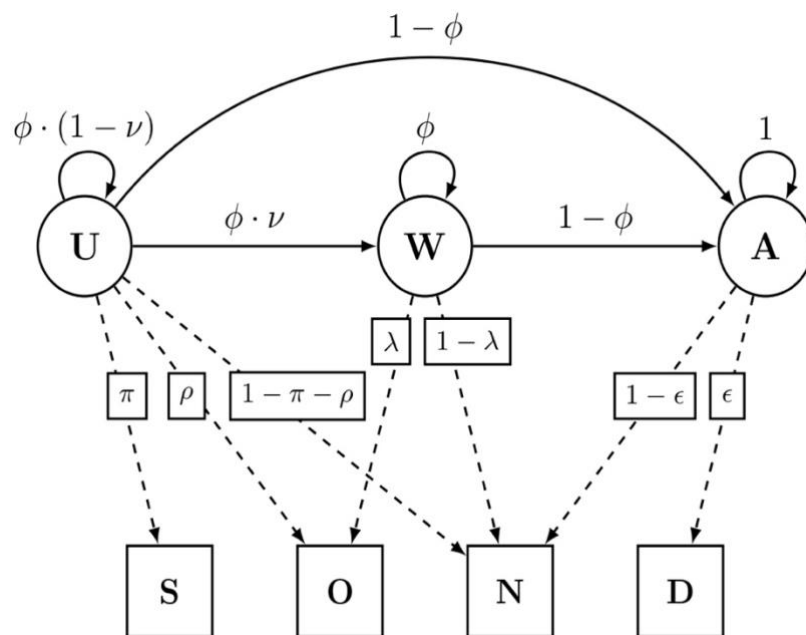
164

165 Throughout the study, we define pups as individuals younger than 6 months, after which all GSL  
166 are referred to as juveniles until they reach 5 years of age and are assumed to be adults (Trillmich  
167 et al. 2016). Due to high site fidelity and limited exploration, deaths of individuals under the age  
168 of 1 year were accurately recorded either by recovering the body, disappearance from the site of  
169 birth and entire islet for very young individuals (< 3 months), or repeated observations of the  
170 mother without her offspring over the course of a field season. Deaths of individuals older than 1  
171 year are only recorded if a body is recovered and a positive ID obtained. We used correlated ONI  
172 anomaly (NOAA Climate Prediction Centre 2024) and sea surface temperature ([SST] Galápagos  
173 Climatology Database 2024), aggregated and averaged by reproductive year, to represent food  
174 availability (Soto et al. 2004; Mueller et al. 2011).

175

176 ***2.2. Multi-state capture-recapture model***

177 For this study weaning is defined as the point in time when an offspring stops suckling. We  
 178 employed MSCRM which we frame as an HMM (see *supplementary material*) and used the  
 179 package LaMa (2025) in R (version 4.3.3) for building the HMM likelihood. We assumed three  
 180 possible latent states for offspring: unweaned, weaned, and absent (Figure 1). All animals'  
 181 trajectories begin in their reproductive year of birth when they are certainly unweaned. From there,  
 182 they may transition to the weaned state after a certain number of seasons, followed by the absent  
 183 state after additional season(s). Alternatively, they may transition from the unweaned state directly  
 184 to the absent state. We assumed irreversible state transitions such that an offspring, once  
 185 considered weaned, cannot return to unweaned, and once considered absent, will stay absent. A  
 186 potential trajectory with one state per season could be U, U, U, U, W, W, A. As our primary interest  
 187 lies in age at weaning, we included the animal's age in years as a covariate in the state process.  
 188 Finding the appropriate functional form of this covariate was addressed during model selection  
 189 (see sp. *Methods*).  
 190



**Figure 1.** Both processes of the MSCRM for the observations suckling (S), observed (O), not observed (N), dead (D) generated given the states unweaned (U), weaned (W), and absent (A). Solid lines show possible state transitions. Dashed lines indicate possible observations given each state. Lines corresponding to probabilities of zero and index are omitted for visual clarity.

192 To describe the state dynamics over time, we modelled the survival rate  $\phi_t$  for  $t = 1, \dots, T$  and  
193 the weaning rate  $v_t$  as functions of an animal's age. It should be noted that the survival rate refers  
194 only to apparent survival from the age of one year onwards, as the absent state is comprised of  
195 dead animals as well as those who were no longer seen on the islet. Given the latent state, different  
196 observations could have arisen: an unweaned animal can be observed suckling with probability  $\pi$ ,  
197 observed but not suckling with probability  $\rho_t$ , or not observed ( $1 - \pi_t - \rho_t$ ). In the weaned state,  
198 an animal can either be observed (but not suckling) with probability  $\lambda_t$ , or not observed ( $1 - \lambda_t$ ).  
199 Absent animals can either be recovered dead with probability  $\varepsilon_t$  or not observed ( $1 - \varepsilon_t$ ). We  
200 accounted for observation effort by writing  $\pi_t, \rho_t, \lambda_t, \varepsilon_t$  as functions of the number of observation  
201 rounds in each season  $t$ .

202

203 The variables offspring sex, ONI, offspring birth SMI (as a proxy for maternal fitness), birth  
204 interval and maternal age in years (at the time of the birth of her offspring) were investigated and  
205 their effect assessed by comparing models using the specific subset both with and without the  
206 variable included. SST was investigated during early exploration. GSL are known to be entirely  
207 nutritionally dependent on milk until 12 months (Jeglinski et al. 2012; Piedrahita 2015).  
208 Individuals who died before reaching 1 year of age were therefore unable to provide us with  
209 information to estimate the probability of any of the possible state transitions. Although the  
210 MSCRM may have been flexible enough that the inference on weaning would not change, we were  
211 concerned that the flexibility would become absorbed as the model attempted to explain early age  
212 survival. As this study's central focus was to investigate age-at-weaning in GSL, we decided to  
213 exclude all individuals registered as deceased before the age of 1 year ( $n = 156$ ).

214

215 We used the Viterbi algorithm (Zucchini, MacDonald & Langrock 2016, Chapter 5) to obtain the  
216 most likely state sequence for each animal given the data and the model parameters. Model  
217 selection for the functional form of the age covariate and inclusion of other covariates in the state  
218 process was guided by the AIC whilst a visual inspection of results served as a final criterion for  
219 judging the relevance of added complexity.

220

221 ***2.3. Logistic regression model***



222 LRMs were used to investigate the probability of an offspring becoming a supersuckler. We  
223 classed supersucklers as offspring deemed unweaned by the selected MSCRM i.e. after the 95%  
224 weaning threshold, which conforms with standard statistical practices (Wei et al. 2006; Waldmann  
225 2018) and previous work on this behaviour (Childs et al. 2025). Supersucklers are therefore 4 years  
226 of age or older. In order to understand the drivers behind the likelihood of individuals becoming  
227 supersucklers we analysed both time-invariant and time-variant variables. The former consisted of  
228 offspring age, age at the birth of a sibling, sex, SMI, and maternal age at parturition. The latter  
229 were ONI anomaly average and observation effort for each reproductive year. Additionally, we  
230 incorporated the decoded state of the offspring in each season based on the MSCRM. For an  
231 unbiased comparison between supersucklers and non-supersucklers, we only included non-  
232 supersucklers who according to the MSCRM had not transitioned to the absent state at 4 years old.  
233 This resulted in a dataset of 766 individuals: 650 non-supersucklers and 116 supersucklers.

234

235 Based on this data an LRM was fitted including offspring sex and ONI anomaly. A series of LRMs  
236 were then built using subsets of data to investigate whether maternal fitness and experience affects  
237 the probability of their offspring becoming a supersuckler: 1. maternal age and birth, 2. birth SMI,  
238 90-day average SMI, and both SMI measurements, 3. birth interval, and the data for this latter  
239 LRM. To control for any possible external ONI or cohort effects, we included year of birth as a  
240 random effect in all models. The effect of a variable was informed by significance results and BIC,  
241 which also guided model selection and parameter inclusion for all LRMs. Missings were checked  
242 for informativeness by comparing the proportion of supersucklers in the data group with missing  
243 values, with the data group with non-missing values for that variable.

244

245 To understand the potential reproductive cost of mothers supporting a supersuckler we visually  
246 inspected the relationship between time until the birth of the next sibling and whether that sibling  
247 survived to yearling age for both supersucklers and non-supersucklers. We further investigated the  
248 number of female supersucklers who were reproductively active at the time of supersuckling. Here,  
249 reproductively active was defined as females who either gave birth within the same reproductive  
250 year as having been observed supersuckling - and were therefore pregnant at the time - or had  
251 already given birth to an offspring prior to their first observed supersuckling event. Male  
252 reproductive data were not available for this study. To assess whether supersucklers grow slower

253 during early life than their non-supersuckling counterparts we plotted and visually inspected  
254 length, weight and SMI curves of supersucklers and non-supersucklers, separated by sex and fitted  
255 with a spline model.

256

### 257 **3. Results**

#### 258 *3.1. Multi-state capture-recapture model*

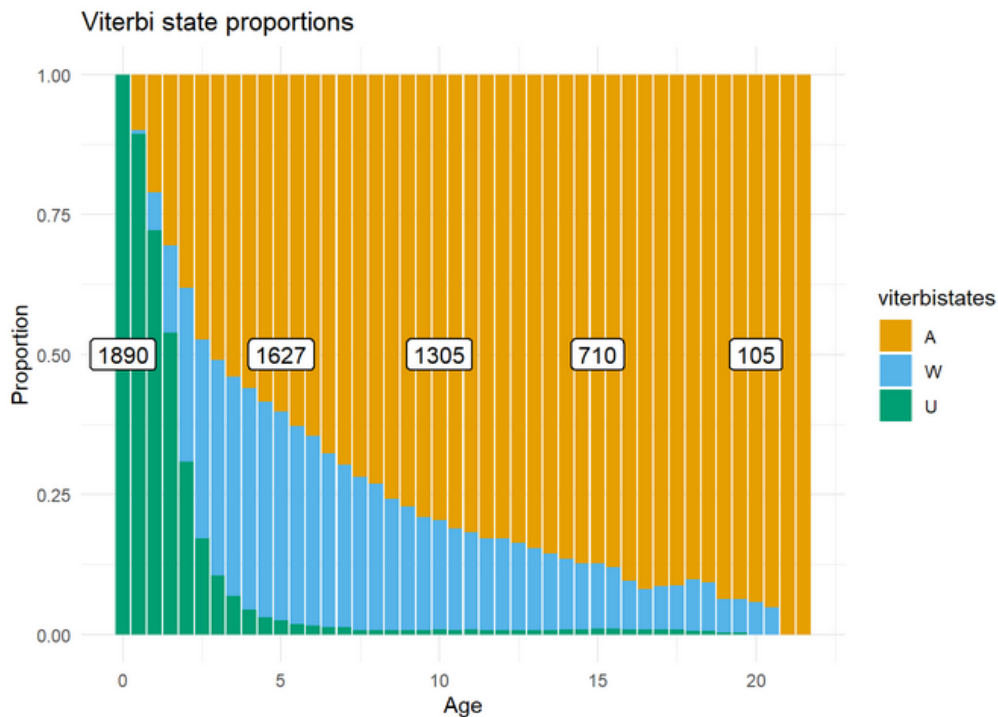
259 All MSCRM model candidates and their fit can be found in Table 1. During early exploration no  
260 effect of SST was found nor was there any improvement to the fit of the MSCRMs, it was therefore  
261 excluded from all models. We found that accounting for observation effort ( $n$  = rounds per year)  
262 increased the model fit. Effort has a direct positive effect on the amount of suckling observed in the  
263 unweaned state: between 56.9% and 86.3% for min. and max. number of observation rounds, as  
264 well as the probability of observing an individual at all (33.1% - 48.2% in the weaned state). Dead  
265 individuals are also more likely to be recovered (0.2% - 2.7% in the absent state) with higher effort.

266 **Table 1.** All fitted MSCRM described by the covariates included in the observation process via Z and the state process via Y. Number of parameters (np) and  
 267 number of individuals (nind) for which the corresponding variables were available are used as a basis for comparison via the AIC. Variable notation  $age_{dec}$  refers  
 268 to the exponential decay function of age  $age^2 * exp^{-age}$ ,  $SMI$  to the SMI of an individual's first 10 days,  $sibl$  to time until sibling,  $m_{age}$  to the age of the mother.  
 269 Index refers to the steps in model selection. The index in bold indicates the model that was preferred at each stage of model selection.

Index	Covariates in Z	covariates in Y	np	nind	AIC
0A	-	-	6	1890	38038.0
<b>0B</b>	<i>rounds</i>	-	10	1890	37586.4
1A	<i>rounds</i>	<i>age</i>	12	1890	37510.8
<b>1B</b>	<i>rounds</i>	<i>age, age<sup>2</sup></i>	14	1890	37091.2
1C	<i>rounds</i>	<i>age, age<sup>2</sup>, age<sup>3</sup></i>	16	1890	37046.4
1D	<i>rounds</i>	<i>age, age<sub>dec</sub></i>	14	1890	37032.0
1E	<i>rounds</i>	<i>I(age<sub>t</sub> = 1), ..., I(age<sub>t</sub> = 14))</i>	38	1890	36821.0
2	<i>rounds</i>	<i>age, age<sub>dec</sub>, sex, sex · age, sex · age<sub>dec</sub></i>	20	1890	37032.2
<b>3</b>	<i>rounds</i>	<i>age, age<sub>dec</sub>, ONI, ONI · age, ONI · age<sub>dec</sub></i>	20	1890	36969.6
4A	<i>rounds</i>	<i>age, age<sub>dec</sub>, ONI, ONI · age, ONI · age<sub>dec</sub>, SMI, SMI · age, SMI · age<sub>dec</sub></i>	26	1266	24529.4
<b>4B</b>	<i>rounds</i>	<i>age, age<sub>dec</sub>, ONI, ONI · age, ONI · age<sub>dec</sub></i>	20	1266	24526.2
<b>5A</b>	<i>rounds</i>	<i>age, age<sub>dec</sub>, ONI, ONI · age, ONI · age<sub>dec</sub>, sibl, sibl · age, sibl · age<sub>dec</sub></i>	26	317	5961.2
5B	<i>rounds</i>	<i>age, age<sup>2</sup>, ONI, ONI · age, ONI · age<sup>2</sup></i>	20	317	5993.4
6A	<i>rounds</i>	<i>age, age<sub>dec</sub>, ONI, ONI · age, ONI · age<sub>dec</sub>, m<sub>age</sub>, m<sub>age</sub> · age, m<sub>age</sub> · age<sub>dec</sub></i>	26	328	no convergence
<b>6B</b>	<i>rounds</i>	<i>age, age<sub>dec</sub>, ONI, ONI · age, ONI · age<sub>dec</sub></i>	20	328	no convergence

271 **Offspring effects**

272 When including age as a functional form (MSCRM 1D), the preferred approach was  
273  $\text{age}^2 \cdot \exp(-\text{age})$ . Although estimating an effect separately for each age category was supported by  
274 AIC, visual inspection showed no meaningful differences to model 1D. As less data was available  
275 for older ages additional complexity was deemed unnecessary. Exponential decay improved the  
276 overall fit compared to simple polynomials without increasing the number of parameters. We  
277 therefore favoured the model in its simplest form, using only exponential decay and the polynomial  
278 effect as parameters. This ensured the flexibility of the polynomial function is prioritised for  
279 younger ages whilst assuming the effect in older ages is captured sufficiently by the linear term.  
280 Viterbi decoding of model 1D provided age-at-weaning estimates of 50% by 1.5 years, 90% by 3  
281 years, and 95% by 4 years (Figure 2). The highest number of state switches was observed between  
282 1 - 2 years of age and rapidly decreased after this point (Figure S1). The highest number of switches  
283 to the weaned state was observed at 1.5 years of age. Switches to the absent state were extremely  
284 elevated throughout the first 2.5 years, peaking at the yearling stage and only levelling out towards  
285 the end of the 3rd year.



**Figure 2.** Viterbi state proportions from Model 1D showing the most likely state (unweaned [U], weaned [W], absent [A]) an individual is in at any given time point given their observational history and age. White-boxed numbers within the figure indicate the total number of possible observable animals at 5-year intervals.

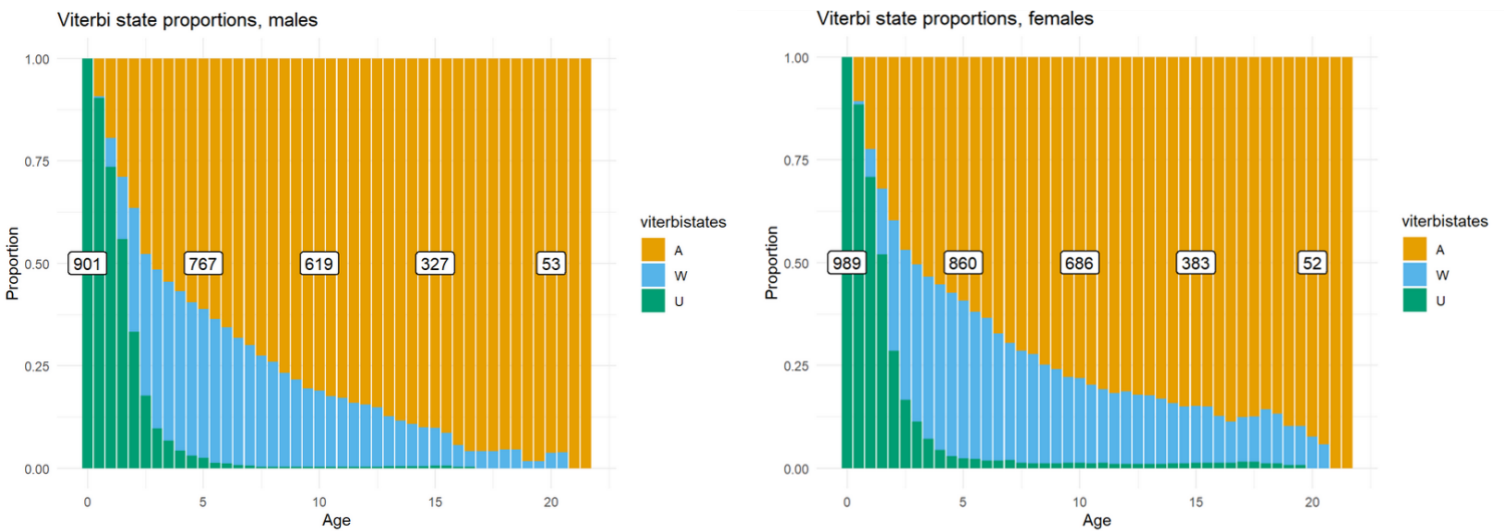
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287 Due to increasing exploration and time spent at sea, the likelihood of recovering a body and  
288 obtaining a positive ID for death registration dwindles dramatically with age. Over the entire data  
289 collection period 247 deaths have been recorded, of which 63% ( $n = 156$ ) were individuals under  
290 1 year of age. Only 40 adult deaths have been recorded. That these numbers are so low is likely due  
291 to most individuals within this population dying at sea or temporary haul-out locations. Without  
292 positive confirmation of these deaths' individuals remain in the dataset even after multiple years of  
293 absence. It should therefore be noted that the sample sizes (Figure 2 & 3) indicate the total number  
294 of possible observable individuals.

295

296 The selection of additional covariates was guided by a combination of AIC, BIC, and contextual  
297 considerations. Among animals that could be observed, both sexes were relatively evenly  
298 represented. We found no apparent effect of sex on weaning age in MSCRM 2, and both visual  
299 inspection and AIC indicated that including sex as a covariate was unnecessary. However, Viterbi  
300 decoding revealed a slight effect on the probability of females transitioning to the absent state later  
301 in life, with males more likely to transition at younger ages (Figure 3).

302



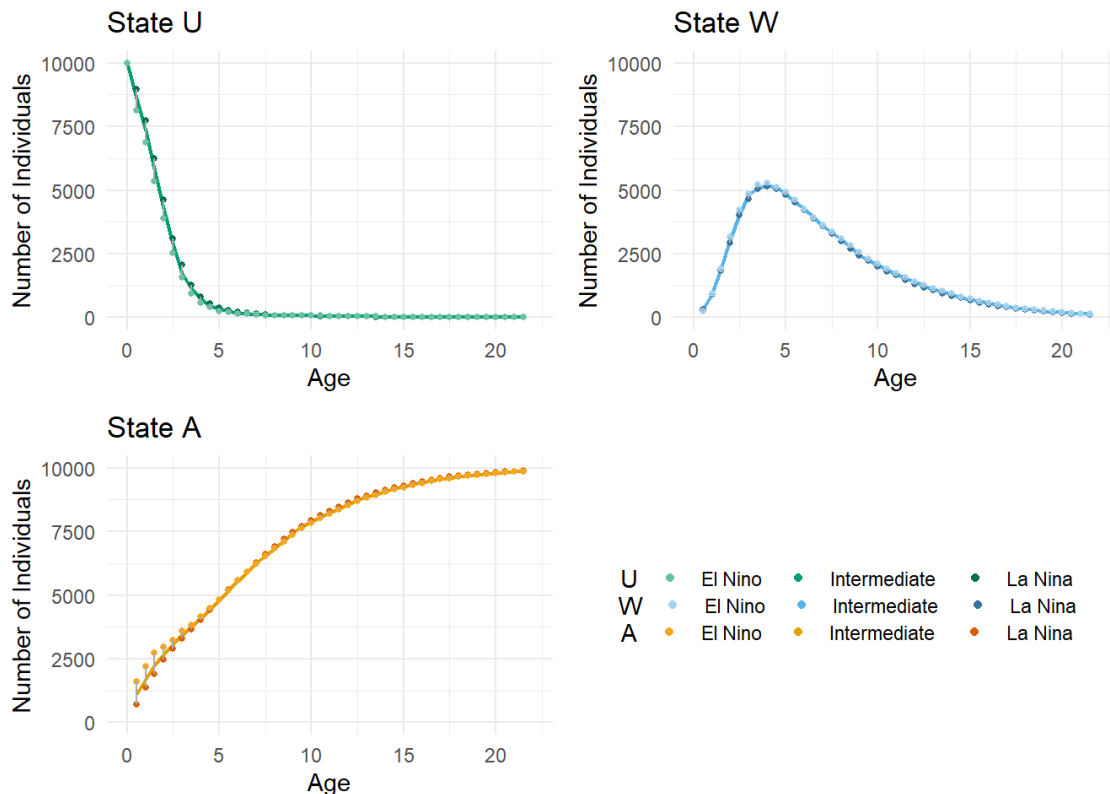
**Figure 3.** Viterbi state proportions from Model 2 showing the most likely state (unweaned [U], weaned [W], absent [A]) of an individual at any given time point is in according to sex. White-boxed numbers within the figure indicate the total number of possible observable animals at 5-year intervals.

303

304 **Environmental effects**

305 Visual inspection and AIC results of MSCRM 3 indicated that ONI should be retained as a  
306 covariate and a visualisation of the relationship (Figure 4) showed that there is a strong effect on  
307 juvenile survival between the ages of 1 - 4 years. This can be seen in the high variation in the  
308 number of individuals in both the unweaned (U) and absent (A) states between El Niño and La  
309 Niña conditions. During poor El Niño conditions individuals are much less likely to be in the  
310 unweaned state and more likely to transition to the absent state, indicating a reduced survival rate  
311 during these events compared to more favourable La Niña periods. No probability on survival prior  
312 to yearling age can be inferred due to the exclusion of these deaths from the model. The probability  
313 of individuals being observed in the weaned (W) state is also affected by ONI between the ages of  
314 2 - 4, where during El Niño individuals between these ages are more likely to be observed in this  
315 state than during La Niña conditions. The effect of ONI on all states reduces with age showing  
316 only a slight influence on weaned and absent between the ages of 7-16.

317



**Figure 4.** ONI category effect on state behaviours unweaned (U), weaned (W) and absent (A). Where the interaction during an Intermediate ONI event is represented by a solid line and El Niño (pale) and La Niña (dark) as points along this line.

318

### 319 **Maternal effects**

320 When investigating maternal life-history effects, information on mothers was unavailable or  
321 incomplete for most offspring, resulting in different subsets when incorporating variables related  
322 to an individual's mother. This provided the base for model fitting.

323

324 Including birth SMI as a covariate is neither supported by AIC nor did the weaning rate in the first  
325 few years of an individual differ given different SMI values. However, SMI did appear to have an  
326 effect on offspring survival (Figure S2) with survival rates estimated between 78% and 96% for  
327 offspring with minimum and maximum SMI values in the data, respectively. When birth interval  
328 was included as a covariate (MSCRM 5A & 5B) AIC results indicated a slight preference for its  
329 inclusion. The MSCRM suggests that a longer interval before the birth of the next sibling is  
330 associated with later weaning (Figure S3) with about 98% of offspring with an early sibling (born  
331 in the same year) weaned by age, but only 61% of offspring with a late sibling (born 7 years later)  
332 based on a simulated state distribution given the model.

333

334 The inclusion of maternal age in MSCRM 6A and 6B produced unstable results, preventing the  
335 identification of a global maximum. This was likely due to the small sample size ( $n = 328$ ) and  
336 specificity of the subset. Due to the colony's decline a large proportion of the females who  
337 recruited and successfully reproduced on Caamaño were first identified as juveniles or adults  
338 during the early years of data collection and therefore have unknown DOB (*personal*  
339 *communication*). Simplification of the models failed to yield more stable results; we were therefore  
340 unable to draw any meaningful conclusions of the effect this life-history trait may have on weaning  
341 age.

342

### 343 **3.2. Logistic regression model**

344 All LRM model candidates and their fit can be found in Table 2. Supersucklers and non-  
345 supersucklers were classified according to the offspring decoded states of MSCRM 3. Due to the  
346 lack of effect of SST in the MSCRM at an annual scale, SST was not included in any LRM. LRM  
347 0B confirmed that sex has no significant effect on the probability of becoming a supersuckler.  
348 Averaged ONI anomalies were included for the year of birth and the first four years of their life as

349 a covariate, in order to understand whether ONI during offspring's early life or the year they  
 350 become a supersuckler are more important. We found that an increase in ONI anomaly during an  
 351 individual's 2<sup>nd</sup> and 3<sup>rd</sup> year of life has a negative impact on their probability of becoming a  
 352 supersuckler. Yet the inclusion of ONI anomalies did not greatly improve the model, with BIC  
 353 indicating that no covariate inclusion was a better fit. In all other LRMs we therefore included the  
 354 year of birth as a random effect to control for any cohort effect, and only ONI anomalies of the 2<sup>nd</sup>  
 355 and 3<sup>rd</sup> year as covariates.

356

357 **Table 2.** All fitted LRMs predicting the probability for an individual to become a supersuckler described by the  
 358 predictors and random effect included. The number of parameters (np) and number of individuals (nind) for which the  
 359 corresponding variables were available are used as a basis for comparison via the BIC. Variable notation  $ONI_0 -$   
 360  $ONI_4$  refers to the oscillation index at the given age of an individual in,  $SMI_{10}, SMI_{90}$  to the SMI of an individual's  
 361 first 10 or 90 days, *sibl* to time until sibling,  $m_{age}$  to the age of the mother. Index refers to the steps in model selection.  
 362 The index in bold indicates the model that was preferred at each stage of model selection.

363

<b>Index</b>	<b>random effect</b>	<b>predictors</b>	<b>BIC</b>	<b>nind</b>	<b>np</b>
<b>0A</b>		-	<b>658.04</b>	766	1
0B		<i>sex, ONI<sub>0</sub>, ONI<sub>1</sub>, ONI<sub>2</sub>, ONI<sub>3</sub>, ONI<sub>4</sub></i>	677.24	766	7
1A	<i>cohort</i>	$m_{age}$	102.91	98	3
<b>1B</b>	<i>cohort</i>	-	<b>98.41</b>	98	2
2A	<i>cohort</i>	$SMI_{90}$	425.74	492	3
2B	<i>cohort</i>	$SMI_{10}, SMI_{90}$	430.38	492	4
2C	<i>cohort</i>	$SMI_{10}$	426.44	492	3
<b>2D</b>	<i>cohort</i>	-	<b>422.27</b>	492	2
3A	<i>cohort</i>	<i>sibl</i>	99.87	140	3
<b>3B</b>	<i>cohort</i>	-	<b>95.97</b>	140	2

364



365 The results of LRM 1A indicate that maternal age at birth does not play an important role in the  
366 probability of offspring becoming a supersuckler. The proportion of offspring without a known-  
367 age mother was similar for supersucklers and non-supersucklers, however, we still assume some  
368 structure in the missingness that might not have been addressed in this model setup. Within this  
369 LRM only a few offspring with older mothers (+12 years) were included; since the data collection  
370 period did not span more than a single lifetime, the younger the mother, the more likely her age  
371 was to be in the data set. We therefore concluded that if there was an effect, we might not be able  
372 to find it, and that this variable should be excluded from further analysis.

373

374 We found no relationship between becoming a supersuckler and SMI at birth or early life SMI and  
375 so excluded this variable from further analysis. It should be noted that the proportion of missing  
376 values for SMI was slightly higher for supersucklers than non-supersucklers, likely a reflection of  
377 colony proportional differences. There was also no indication of a relationship between birth  
378 interval to the next sibling and the probability of an offspring becoming a supersuckler. Due to the  
379 exclusion of mothers with recorded absence (2+ years) from the database and the previously stated  
380 data imbalance, the proportion of missings for supersucklers was again much higher than for non-  
381 supersucklers. Additional visualisation of birth interval and survival of the next sibling to yearling  
382 age (Figure S4), suggested that there may be some relationship between sibling first-year survival  
383 and older offspring becoming a supersuckler but further highlighted the high number of missing  
384 values. We concluded that, due to the small and imbalanced sample size, it was unlikely that we  
385 might be able to find the effect if there was one and that this variable could be excluded from  
386 further analysis.

387

388 Visual inspection of weight, length and SMI curves (Figure S5) showed high overlap between the  
389 predicted means for pups who would become supersucklers and those who would not, indicating  
390 no difference in growth between supersucklers and non-supersucklers. The widening of the  
391 confidence intervals (95%) from 700 days (1.9 years) onwards indicates that the curve becomes  
392 implausible at this point due to the reduced sample size.

393

#### 394 **Conflict of Interest**

395 None declared.

396

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406

407 **Supplementary material**

408

409 **1. Methods: Model formulation of the MSCRM**

410 The multi-state capture-recapture model (MSCRM) belongs to the class of hidden Markov models  
411 (HMMs) comprising two stochastic processes: the state process and the observation process.  
412 HMMs assume that (partially) hidden states determine the distribution of the observations at each  
413 point in time.

414

415 State Process:

416 The state process  $\{S_t\}_{t=1,\dots,T}$  is defined as a Markov chain with  $s_t$  the state at season  $t$ . It aims to  
417 describe the weaning process by distinguishing the unweaned state "U" and the weaned state "W".  
418 Additionally, the absent state "A" accounts for events of death and emigration. This absorbing  
419 state ensures that dead or absent animals which cannot be observed anymore are not considered  
420 weaned and the weaning process is therefore only informed by animals that can be observed. The  
421 upper half of the directed graph in Figure 1 shows the possible state transitions. Notably, once the  
422 unweaned state is left, an animal cannot transition back to it. The same applies for the weaned  
423 state. Therefore, the following transition probability matrix  $\Gamma =$

424  $(\gamma_{UU} \ \gamma_{UW} \ \gamma_{UA} \ 0 \ \gamma_{WW} \ \gamma_{WA} \ 0 \ 0 \ 1) = (\phi \cdot (1 - \nu) \ \phi \cdot \nu \ 1 - \phi \ 0 \ \phi \ 1 - \phi \ 0 \ 0 \ 1)$

425 is assumed with  $\gamma_{kl}$  denoting the probability of switching from state  $k$  to  $l$  for all  $k, l \in \{U, W, A\}$ .

426

427 In defining a survival rate that reflects the probability that an animal continues to be observable,  
428 and the assumption that survival rate is independent of the current state, we could then rewrite the  
429 entries in  $\Gamma$  in terms of this survival rate  $\phi$  and a weaning rate  $\nu$ . In practical terms, by estimating  
430 only  $\gamma_{UU}$  and  $\gamma_{UW}$  and considering that  $\gamma_{UU} + \gamma_{UW} + \gamma_{UA} = 1$ , we can derive  $\phi = \gamma_{UU} + \gamma_{UW}$

431 and  $\nu = \frac{\gamma_{UU}}{\gamma_{UU} + \gamma_{UW}}$ .

432

433 Covariates that were assumed to affect the weaning and survival rate  $(\nu, \phi)$  could be included in  
434 the state process leading to an inhomogeneous Markov chain  $\{S_t\}$  with  $\Gamma_t$  and therefore  $(\nu_t, \phi_t)$   
435 varying over time. In order to do so, the transition probabilities to be estimated are linked to linear  
436 predictors  $\tau_{j,t} = Y_t \beta_j$  for  $j = 1, 2$  and  $t = 1, \dots, T \#(1)$  given the covariate matrix  $Y_t$  and the

437 regression coefficients  $\beta_j$  such that  $\gamma_{UU,t} = \frac{\exp(\tau_{1,t})}{1 + \exp(\tau_{1,t}) + \exp(\tau_{2,t})}$  and  $\gamma_{UW,t} =$   
 438  $\frac{\exp(\tau_{2,t})}{1 + \exp(\tau_{1,t}) + \exp(\tau_{2,t})}$ .

439

440 Observation process:

441 The observation process  $\{X_t\}_{t=1,\dots,T}$  depends on the underlying state process  $\{S_t\}$  as depicted in  
 442 the bottom half of Figure 1. The observations  $x_{t,q}$  correspond to being observed suckling (S), being  
 443 observed but not suckling (O), not being observed (N), or recovered dead (D) for each field season  
 444  $t$  and each animal  $q$ .

445

446 We constructed three state-dependent probability distributions as follows:

447  $X|j \sim \text{Categorical}(q_{\cdot|j})$ , for  $j = U, W, A$

448 
$$q_{\cdot|U} = (\pi, \rho, 1 - \pi - \rho)$$

449 
$$q_{\cdot|W} = (0, \lambda, 1 - \lambda, 0)$$

450 
$$q_{\cdot|A} = (0, 0, 1 - \epsilon, \epsilon)$$

451

452 Given the unweaned state  $U$ , we obtained a categorical distribution where individuals who were  
 453 observed suckling had probability  $\pi$ , observed doing something else had probability  $\rho$  and  
 454 unobserved with probability  $1 - \pi - \rho$ . Given the weaned state  $W$  and the absent state  $A$ , the  
 455 distributions are essentially Bernoulli with  $\lambda$  the probability of being observed given the weaned  
 456 state and  $\epsilon$  the probability of being recovered dead.

457

458 We collect variables that are assumed to affect the resighting probability and thereby the  
 459 distribution of observations in a covariate matrix  $Z_t$  and build a linear  $\eta_{i,t} = Z_t \alpha_i$  for  
 460  $i = 1, 2, 3, 4$  and  $t = 1, \dots, T$  with  $\alpha_{i,t}$  the regression coefficients.

461

462 The four parameters  $\rho_t, \pi_t, \lambda_t, \epsilon_t$  are then time varying by being linked to the linear predictors as  
 463 follows:

464 
$$\rho_t = \frac{\exp(\eta_{1,t})}{1 + \exp(\eta_{1,t}) + \exp(\eta_{2,t})}$$

465

$$\pi_t = \frac{\exp \exp (\eta_{2,t})}{1 + \exp \exp (\eta_{1,t}) + \exp \exp (\eta_{2,t})}$$

466

$$\lambda_t = \frac{1}{1 + \exp \exp (\eta_{3,t})}$$

467

$$\epsilon_t = \frac{1}{1 + \exp \exp (\eta_{4,t})}$$

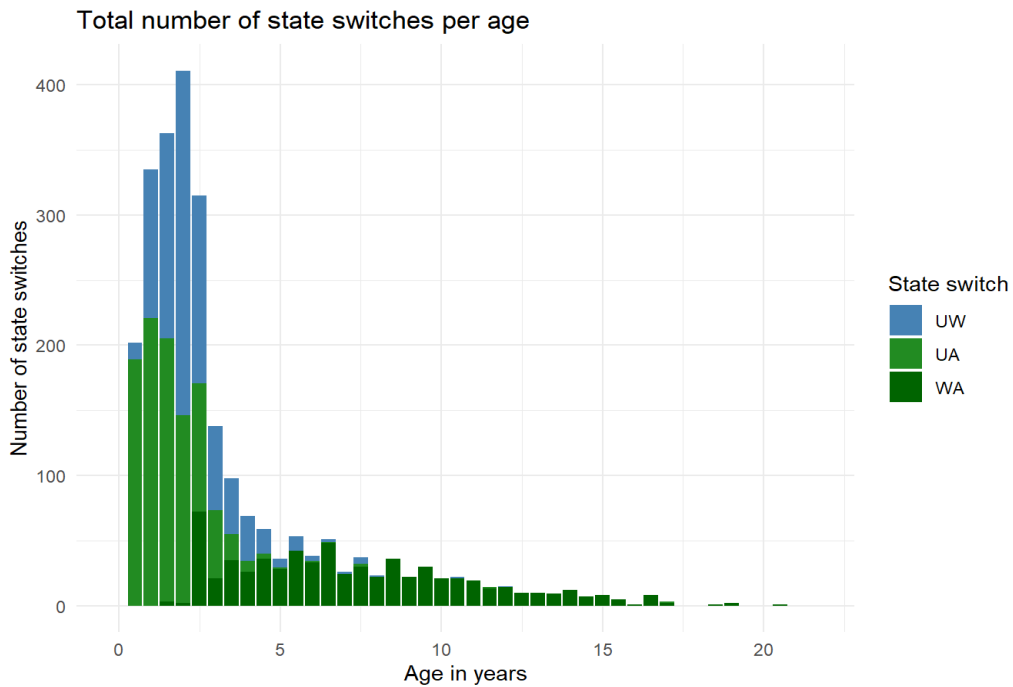
468

469 Parameter estimation and state decoding:

470 By framing the model as a hidden Markov model, we leveraged inferential tools like the forward  
 471 algorithm to estimate the latent weaning process and the Viterbi algorithm to decode the most  
 472 likely state sequence. Note that we constructed a partially hidden Markov model in the sense that  
 473 a suckling observation "S" and a death recovery "D" inform directly about the state having  
 474 generated this observation. We used the package LaMa in R for conveniently building the HMM  
 475 likelihood and maximized the likelihoods directly by using the numerical optimizer nlm.

476

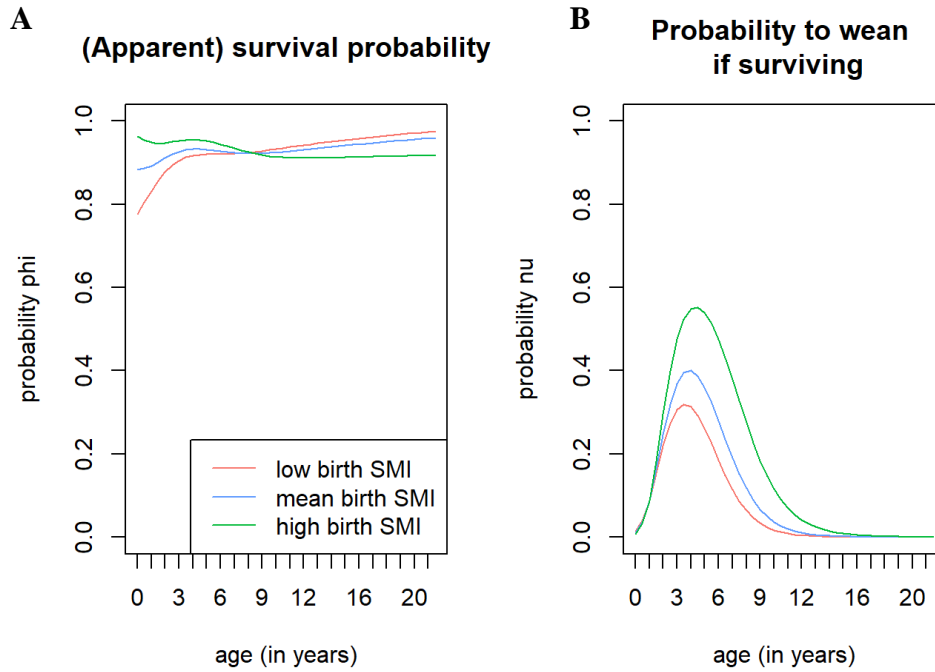
477 **2. Results**



**Figure S1.** Total number of switches between each state according to offspring age: unweaned-weaned (UW, blue), unweaned-absent (UA, light green), weaned-absent (WA, dark green).

478

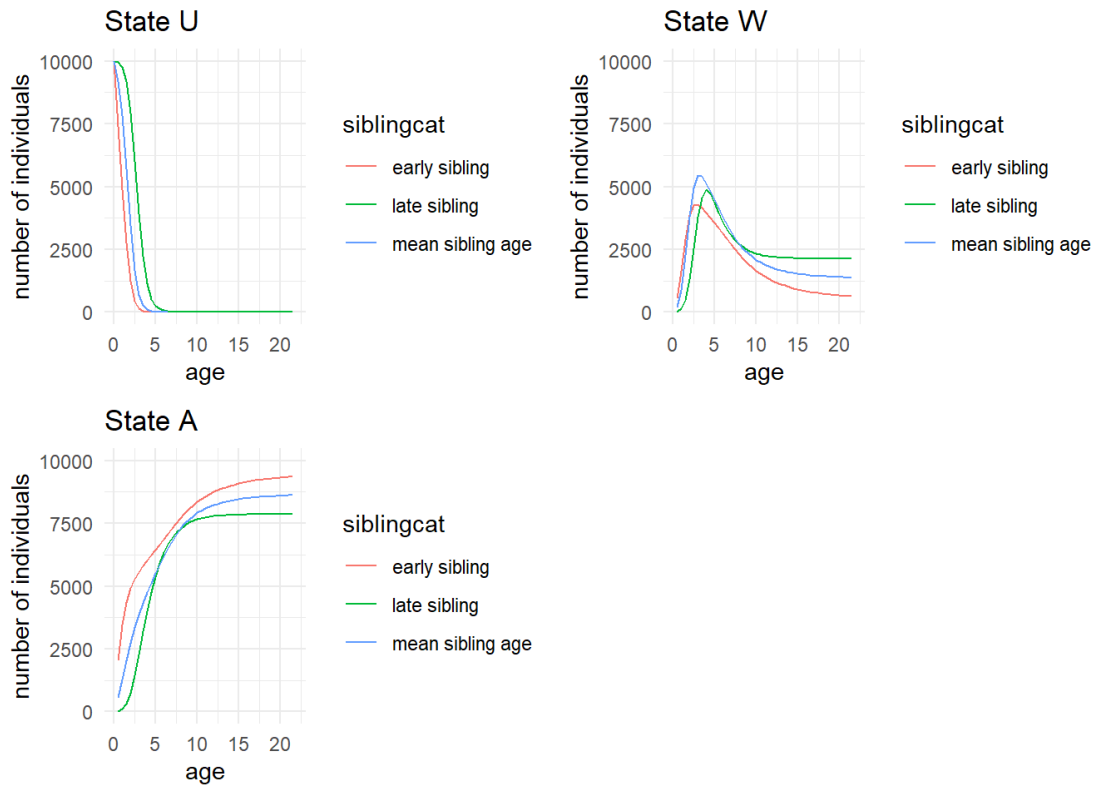
479



**Figure S2.** Apparent survival probability (A) of offspring according to their birth SMI, indicated by red (low birth SMI), blue (mean birth SMI) and green (high birth SMI) lines, and their age. As well as their probable age at weaning if they survive (B).

480

481



**Figure S3.** Birth interval effect - calculated as time until the birth of the next sibling - on state behaviours unweaned (U), weaned (W) and absent (A) according to offspring age. Indicated by solid lines where red is a short interval (early sibling = 1 year), blue is a mean interval and green a long interval (late sibling = 7 years) they survive (B).





487

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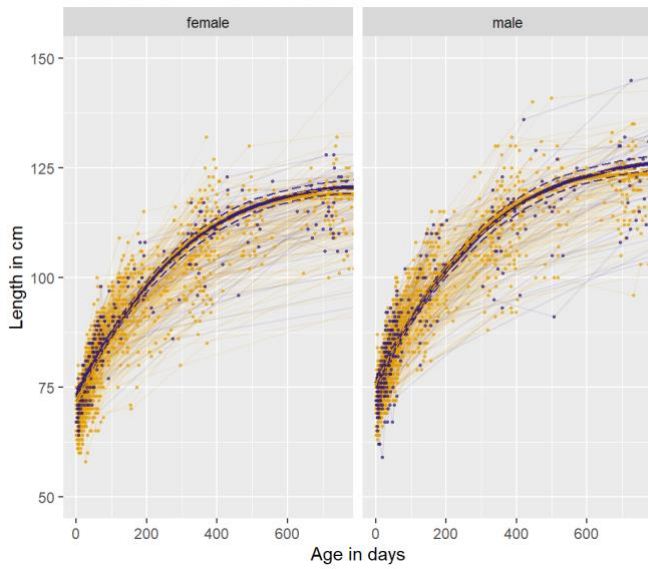
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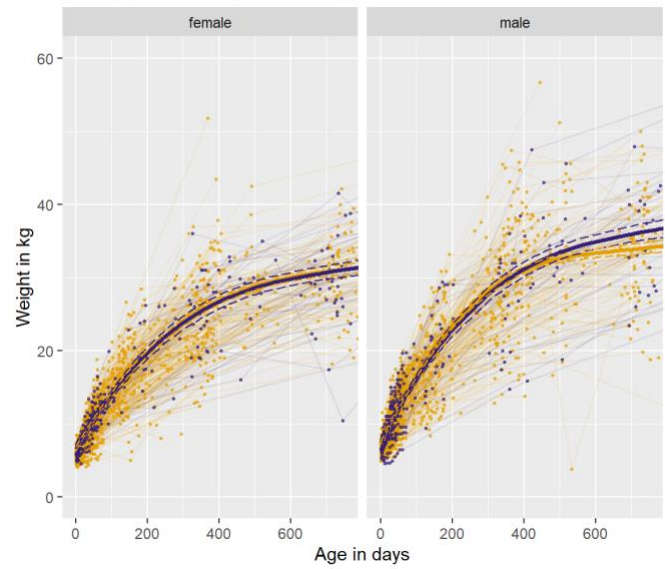
491

492

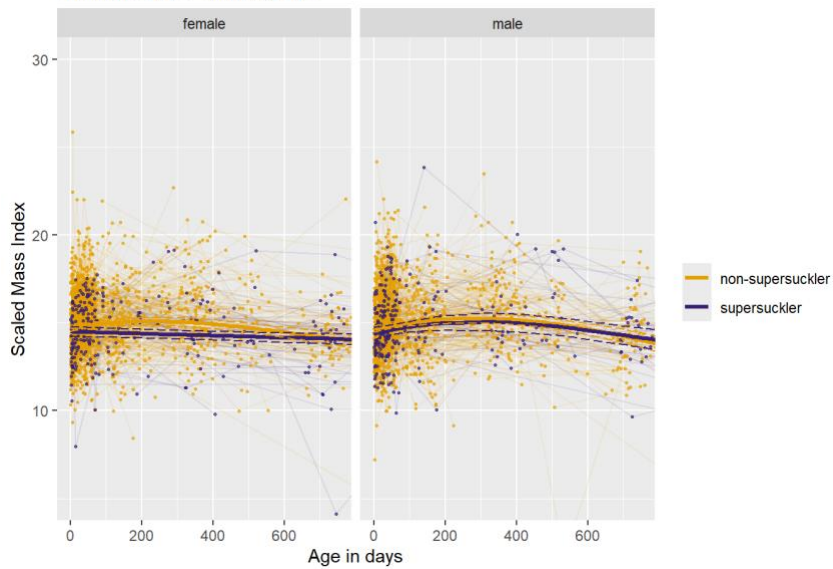
**A** Growth curve: length, 2 years



**B** Growth curve: weight, 2 years



**C** Growth curve: SMI, 2 years



**Figure S5.** The thick lines show the predicted average **a)** body weight, **b)** length and **c)** SMI given the age and sex for individuals later becoming supersucklers (orange) and non-supersucklers (purple). The dashed lines indicate the 95% confidence intervals of the predicted mean.

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