

Lifespan and reproductive cost explain interspecific variation in the optimal onset of reproduction

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110 Fitness can be profoundly influenced by the age at first reproduction (AFR), but to date the 111 AFR-fitness relationship only has been investigated intraspecifically. Here we investigated 112 the relationship between AFR and average lifetime reproductive success (LRS) across 34 bird 113 species. We assessed differences in the deviation of the Optimal AFR (i.e., the species-114 specific AFR associated with the highest LRS) from the age at sexual maturity, considering 115 potential effects of life-history as well as social and ecological factors. Most individuals 116 adopted the species-specific Optimal AFR and both the mean and Optimal AFR of species 117 correlated positively with lifespan. Interspecific deviations of the Optimal AFR were 118 associated with indices reflecting a change in LRS or survival as a function of AFR: a delayed 119 AFR was beneficial in species where early AFR was associated with a decrease in subsequent 120 survival or reproductive output. Overall, our results suggest that a delayed onset of 121 reproduction beyond maturity is an optimal strategy explained by a long lifespan and costs 122 of early reproduction. By providing the first empirical confirmations of key predictions of 123 life-history theory across species, this study contributes to a better understanding of life-124 history evolution.

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126 KEY WORDS: Age at first reproduction, comparative method, cost of reproduction, family

- 127 formation theory, life-history theory.
- 128 **DATA ARCHIVING:**
- 129 Data are provided in the appendix.
- 130 ABBREVIATIONS: AFR, age at first reproduction; LRS, lifetime reproductive success; LRT,

131 likelihood ratio test

132 Life-history theory predicts that the timing of reproductive events during an individual's life 133 affects its fitness (Cole 1954; Caswell 1982). An early age at first reproduction (hereafter 134 AFR) can increase the number of lifetime reproductive events and shorten generation time, 135 which, in a stable or growing population, should be favored by natural selection (Cole 1954; 136 Bell 1980; Roff 1992; Charlesworth 1994). However, an early AFR may also be costly and 137 reduce future survival or reproductive investment (Lack 1968; Roff 1992; Stearns 1992). 138 Additionally, individuals could benefit from deferring breeding beyond sexual maturity if this 139 enhances parenting skills ('constraint hypothesis': Curio 1983), secures access to higher 140 quality territories or mates ('queuing hypothesis': Zack and Stutchbury 1992; van de Pol et 141 al. 2007), increases reproductive output with age ('restraint hypothesis': Williams 1966; 142 Forslund and Pärt 1995) or decreases reproductive senescence ('senescence hypothesis': 143 Charmantier et al. 2006). If AFR is shaped by natural selection, then individuals should adopt 144 the AFR that is associated with the highest fitness return, which may depend on individual 145 guality and annual variation in environmental conditions.

146 Individuals of some species express no variation in AFR, while there is a large range 147 in AFR in other species. In the latter case, only certain AFRs are associated with a high 148 lifetime reproductive success (hereafter LRS), but the exact association appears to vary 149 among species (Clutton-Brock 1988; Newton 1989; Oli et al. 2002; Krüger 2005; Charmantier 150 et al. 2006; Millon et al. 2010; Kim et al. 2011; Tettamanti et al. 2012; Zhang et al. 2015). 151 Moreover, the relationship between the species-specific AFR that is associated with the 152 highest LRS (hereafter termed Optimal AFR) and age of sexual maturity can vary across 153 species (Komdeur 1996; Pyle et al. 1997; Oli et al. 2002; Krüger 2005). Yet, the reasons 154 underlying this among-species variation remain unclear as we currently lack comparative 155 studies that investigate the evolution of AFR and deviation in the timing of Optimal AFR

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during reproductive lifespan across species. Such a study could contribute to our 157 understanding of the general patterns of variation in this crucial life history trait.

158 Whether species-specific Optimal AFR either approximates or is shifted beyond the 159 age of sexual maturity of the species may depend on interspecific variation in life-history or 160 ecological factors. Across species, the pace of life (i.e. slow or fast life history) is likely to be 161 a major factor influencing variation in AFR and timing of the species-specific Optimal AFR 162 relative to the age of sexual maturity (Roff 1992; Stearns 1992; Charlesworth 1994). A short 163 lifespan should be associated with little or no variation in AFR, and with an Optimal AFR that 164 is close to the species' age of maturity, as any postponement would increase the risk of 165 death before reproduction. In contrast, a long lifespan allows for a larger range in AFR and 166 increases the likelihood of a delayed Optimal AFR, an outcome that is supported by field 167 studies (Pyle et al. 1997; Tettamanti et al. 2012). In addition to lifespan, other life-history, 168 ecological or social traits may influence the deviation from the age of sexual maturity in the 169 species-specific Optimal AFR. Species could benefit from delayed AFR when there is a high 170 level of parental care (e.g. altricial species), or when requiring time to learn specialized skills 171 to survive or reproduce successfully. Conversely, a prolonged association of juveniles with 172 their parents (i.e. family-living; Drobniak et al. 2015) may facilitate skill learning and lead to 173 an earlier species-specific Optimal AFR ('skill hypothesis': Skutch 1961; Langen 1996). An earlier Optimal AFR may also be found in cooperatively-breeding species, since helpers may 174 buffer the reproductive costs of early AFR ('load-lightening hypothesis': Khan and Walters 175 176 2002; Santos and Macedo 2011).

177 Here, we use data from 34 bird species to investigate the extent of variation in 178 reproductive strategies and to assess the potential benefits some species may gain from 179 delaying AFR beyond sexual maturity. We examine interspecific variation in the fitness

180 consequences of AFR using within-species relationships between AFR and a fitness proxy 181 averaged over all individuals within a specific AFR-class. For each of the 34 species, we 182 identified the species- and sex-specific Optimal AFR and several derived metrics, 183 summarized in Table 1, to assess changes in LRS or survival as a function of AFR. Information 184 on species-specific Optimal AFR was previously unavailable for typical meta-analysis approaches due to the substantial challenge of obtaining fitness estimates of populations 185 186 from several species. Its investigation allows us to make inferences about the selection 187 pressures on AFR that could not be achieved via a simple analysis of interspecific variation in 188 AFR. As a fitness proxy, we used the most commonly provided measure of an individual's 189 productivity, the lifetime number of fledglings or recruits produced (LRS) (Clutton-Brock 190 1988; Newton 1989 and other references in Table S1). Although it depends on population 191 dynamics, while rate-sensitive fitness estimates (e.g. lambda λ_{ind}) theoretically are more 192 accurate proxies than LRS (Cole 1954; Lewontin 1965; Caswell and Hastings 1980), a number 193 of studies have shown that LRS is a reliable estimate of fitness (Brommer et al. 2002; Link et 194 al. 2002; Dugdale et al. 2010).

195 Specifically, we addressed the following three questions: (i) How does AFR vary 196 within and among species? (ii) Is variation in AFR associated with differences in LRS, and is 197 the typical AFR of a species the one associated with the highest LRS? (iii) Which life-history 198 (chick developmental mode, LRS and survival change with AFR, lifespan), social (family-199 living, helper presence) and ecological (latitude, nest predation) factors are associated with 200 among-species variation in deviation of the Optimal AFR from age at maturity? We used a 201 generalized linear mixed model approach in a model selection framework for the analyses, 202 with further control for similarity in phenotype among taxa due to a shared phylogenetic 203 history.

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205 *Materials and Methods*

206 DATA COLLECTION

207 We used data from published (N = 15) and unpublished (N = 21) studies on the age at first 208 reproduction (AFR) and lifetime reproductive success (LRS) for 34 avian species (Table S1). 209 To find published data, we searched online databases (ISI Web of Science, Scopus) using the 210 terms "age at first reproduction", "age at first breeding", or "age at maturity" in 211 combination with "lifetime reproductive success", "lifetime reproductive output", or 212 "fitness" and "avian" or "bird". We included data from long-term studies (years of 213 monitoring exceeding the mean lifespan) in which individuals were followed for a sufficient 214 period to accurately measure LRS (mean duration of study: 20.75 years; range: 8 to 48 215 years) and where LRS (including its mean, standard deviation and sample size) was reported 216 separately for each category of AFR. We used GetData Graph Digitizer 2.25 217 (http://www.getdata-graph-digitizer.com/) to extract values from published data that were only presented in figures. Unpublished data were requested from researchers who 218 219 coordinated long-term monitoring studies.

220 We collected species-specific data on key life-history, ecological and social lifestyle 221 factors that might influence the effect of AFR on LRS (italicized words represents variable 222 names used in the models), including chick development mode (altricial or precocial), mean 223 lifespan, mean body mass, latitude, nest predation risk, family-living and helper presence. 224 We also collected data on the age of maturity for the estimation of an index used as 225 variables in the model (see INDICES AND ESTIMATES). Age of maturity corresponded to the 226 age at which an individual is physiologically able to reproduce, or the minimum age 227 recorded for breeders. Among ecological factors that can contribute to nest predation risk,

228 nest location is well known and important (Martin and Li 1992; Martin 1993). Based on this 229 information, we ordinally ranked the nest predation risk as high risk – ground nesters, 230 medium risk – nests in shrubs, low risk – nests in trees, or very low risk – cavity breeders or 231 species that build their nest floating on water and thus difficult for nest predators to access. 232 We considered species to be family-living when offspring remain with the parents beyond 233 independence and non-family living when juveniles disperse soon after becoming 234 independent (Drobniak et al. 2015). Species were categorized with helper when offspring 235 regularly engage in cooperative breeding and without helper when offspring do not engage 236 in cooperative breeding. Variables not provided for the populations studied were obtained 237 from the Animal Ageing and Longevity database (http://genomics.senescence.info/species/) or the Handbooks of the Birds of the World (del Hoyo et al. 1992-2006). 238

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240 DATA COMPOSITION

The 34 species included in our study (Figure S1) comprise 10 taxonomic orders and 22 241 242 families, with mean lifespan ranging from 1.4 to 18.5 years and mean LRS ranging from 0.67 243 to 21.16 fledglings produced over the lifetime, or from 0.54 to 2.53 recruits. For blue tits 244 (Cyanistes caeruleus) and western gulls (Larus occidentalis), we included data from two 245 different populations that were analyzed separately. While age at first reproduction might be influenced by individual quality (Forslund and Pärt 1995; Kim et al. 2011), only few 246 247 studies provide such information, limiting our ability to include this factor in our analyses. 248 Data collected consisted of average values per species (i.e. body mass) or per AFR age-class 249 category combining data from all cohorts and years. Therefore, annual or cohort variation 250 could not be addressed here but we hope to do so in future work. Note that not controlling 251 for intraspecific individual quality and combining data across cohorts and years is

252 conservative as it reduces the chance of observing biological patterns. Values of mean LRS 253 (N = 34 species) and lifespan (N = 21 species), as well as their standard deviation and sample 254 size (number of individuals), were determined for each AFR age-class category (e.g. from all 255 individuals starting to reproduce at AFR = 1-year old, at AFR = 2, at AFR = 3, and so on), and 256 for each sex if possible. While it would have been more appropriate to use the geometric rather than the arithmetic mean, as it takes into account variability in fitness (see Liou et al. 257 258 1993), such data were unavailable. Age at first reproduction was defined as the age at which 259 an individual first reproduced during its life. In most species, this value corresponds to the 260 age when a female laid at least one egg, although in some species the value reflects when a 261 female laid a full clutch. For males, AFR corresponds to the age where its mate laid eggs, 262 and, accordingly, reproductively competent males that failed to acquire a mate were not 263 considered as reproductive at that time. The LRS data were based on the number of 264 fledglings or recruits produced over the lifetime of an individual (Table S1). All LRS values 265 were centered and scaled within species and sexes to convert the original units to those of standard deviations and make them comparable (Schielzeth 2010). For species with only 266 267 one AFR age-class category, only a single data point was available. Thus we could not 268 estimate the standard deviation necessary for scaling. Instead, we used the standard 269 deviation of the same sex of a species with a similar value of unscaled LRS to calculate the 270 scaled LRS. Accurate estimation of AFR and fitness proxies is challenging as it requires 271 known-aged individuals and intensive individual-based monitoring of reproductive output 272 throughout the lifespan of a representative sample of individuals, as well as data on the 273 survival and reproduction of descendants. Age at first reproduction and fitness proxies may 274 be biased due to extra-pair paternity, or because not all reproductive events of individuals 275 are followed due to emigration from or immigration into the study population.

276 Consequently, AFR might be overestimated and LRS underestimated for males and 277 overestimated for females. Such biases affect the interpretation of the relationship between 278 AFR and fitness components, and add noise to the data. However, because a relation 279 between AFR and extra-pair paternity and or migration has never been documented, we do 280 not know how and to what extent such a bias would affect our interpretation.

281

282 INDICES AND ESTIMATES

283 Interspecific variation in deviations of the Optimal AFR from the age at sexual maturity 284 might be explained by the association of an early or a late AFR with an increase or a 285 decrease in subsequent survival or reproductive output. However, given the heterogeneity 286 of the data distribution between species and sexes, conventional methods are unable to 287 estimate changes in reproductive output or survival with a changing AFR. Thus, we 288 calculated five derived metrics from the raw data per AFR age-class category to investigate 289 this hypothesis (i.e. average values over all individuals from a specific AFR age-class, 290 combining cohorts and years, for each species and where possible split by sex). These 291 included the Delay Index, which assessed the deviations of the Optimal AFR from the age at 292 sexual maturity, and four indices which assess the relationship between AFR and LRS or 293 survival: the Before Variation Index and the After Variation Index, the Choice Index, and the 294 Lifespan Effect Index (see Table 1).

We visually determined the species-specific AFR that maximized LRS ("Optimal AFR"-Table 1). The use of a single statistical optimization method was not feasible due to the large diversity of patterns in the relationship between AFR and LRS.

Based on the Optimal AFR, the age at sexual maturity and the latest AFR observed within focal species and sex, we assessed the "Delay Index" representing the timing of the Optimal AFR in relation to the reproductive lifespan (illustrated in Table 1):

$$Delay Index = \frac{Optimal AFR - maturity age}{latest AFR - maturity age}$$

301 A Delay Index equal to zero always resulted from the Optimal AFR being the age of maturity. 302 For 35 out of 62 cases several AFR categories had mean LRS values near that of the 303 Optimal AFR. Hence, we determined the range of the species-specific optimum ages for the 304 onset of reproduction, referred as the "Optimal AFR Range". The Optimal AFR Range 305 included the AFR categories adjacent to the Optimal AFR, with mean LRS values included in 306 the calculation of the standard error bar for the mean LRS of the Optimal AFR (Table 1). The 307 AFR categories forming the Optimal AFR Range are therefore assumed to be similarly 308 beneficial in terms of LRS than the Optimal AFR.

309 Based on the Optimal AFR Range, we estimated the Before Variation Index and the 310 After Variation Index. These indices correspond to the slope of the relationship between LRS 311 and AFR from the earliest and the latest AFR to the center of the Optimal AFR Range. The 312 slopes were estimated in the whole data set with all AFR age-class categories, and in a data 313 set only including categories with more than 5% or 10% of the individuals (Table 1). Before 314 and After Variation Indices represent the average of the three estimated slopes. We 315 assumed that a delayed AFR should be favored if an early AFR is associated with a lower LRS, 316 while an earlier AFR should be favored if a late AFR is associated with a lower LRS. Therefore, we expected the Delay Index to be positively correlated with the Before Variation 317 318 Index but negatively with the After Variation Index.

Based on the Optimal AFR Range and the actual value observed for the AFR, we calculated the Choice Index (Table 1), which represented the probability that individuals adopt AFR(s) with highest fitness return:

Choice Index =
$$\frac{\text{Optimal AFR Range}}{\text{number of AFR categories}}$$

322 In cases with only one AFR category (N = 6 out of 62 cases), the Choice Index was assigned a zero, as in such cases there is no variation in AFR. We assumed that species with a large 323 324 Optimal AFR Range relative to the number of AFR categories (i.e. with a large Choice Index) 325 would have a lower probability of suffering a LRS cost when initiating reproduction earlier or 326 later than the Optimal AFR. Consequently, such species may have a higher likelihood of 327 benefiting from delayed reproduction than species with only a low number of beneficial 328 AFR. Therefore, we expected the Delay Index to be positively correlated with the Choice 329 Index.

330 The association between AFR and subsequent survival was calculated via the 331 Lifespan Effect Index, i.e. the correlation coefficient of the reproductive lifespan plotted 332 against AFR per age-class category. We were able to estimate the Lifespan Effect Index for 333 21 out of 34 species only, due to missing data for mean lifespan for the different AFR age-334 class categories for 13 species. As causes and consequences cannot be disentangled from a 335 correlation, negative values could indicate a reproductive cost in terms of survival for 336 individuals with a late AFR or an early AFR favored by high intrinsic mortality. By contrast, 337 positive values could indicate a survival cost of early AFR or a late AFR favored by low intrinsic mortality (Table 1, Figure S2). We assumed a survival cost of early AFR to be 338 339 associated with a late Optimal AFR. Therefore, we expected the Delay Index to be positively 340 correlated with the Lifespan Effect Index.

341 We verified the robustness of our results based on the indices involving the Optimal 342 AFR Range by considering a second method to estimate it. In this second method, the 343 Optimal AFR Range included AFR(s) adjacent to the Optimal AFR with their 90% Cls 344 overlapping those of the Optimal AFR. The first method (method used in the manuscript 345 abovementioned) represents the logic of a null-hypothesis-like test, which assumes an error 346 distribution around the hypothesis (the Optimal AFR's LRS mean), and if our statistics (the 347 other AFRs' LRS mean) do or do not fall within this range. We also considered this first 348 method to be more straightforward while the use of the second method is more 349 conservative. This is because the use of 90% CI indicates that the LRS population's mean of 350 the focal AFR will fail in 90% of the time, while for the use of the standard error it would do 351 so in around 68% of the time. However, we preferred to present the results from the first 352 method in the manuscript for two reasons. First, most of our data comes from studies with 353 intensive monitoring of a population (Table 1, some of which pretty much sample all 354 individuals in the population) and thus, the LRS means approach the population mean with little error. Second, for some AFRs the LRS estimates were based on a single individual (thus 355 356 without CI). Note that one could prefer to consider one or the other method depending on 357 their data characteristics and questions.

358

359 STATISTICAL ANALYSIS

360 General procedure

All statistical analyses were carried out in R version 3.0.2 ((R Core Team 2013), http://www.R-project.org/) using linear mixed-effects models (Imer function, Ime4 package: Bates et al. 2014) that allow for the non-independence of data from a single species by including species as a random factor in the model. To account for differences in sample size

365 (N, Table S1) and decrease noise by giving greater emphasis to the more reliable species-366 specific estimates, all models were weighted (Garamszegi and Møller 2011) by incorporating 367 N-1 in the "weights" argument of the Imer function (Hansen and Bartoszek 2012). Note that 368 removing the weighting did not change the results (Table S2 to S7). To compare coefficients, 369 all continuous predictors were centered (around the mean) and scaled (by the standard 370 deviation) before incorporation in the models (Schielzeth 2010), but we present raw data in 371 the figures. Model assumptions of normality and homogeneity of residuals were checked by 372 visually inspecting histograms and qq-plots of the residuals as well as by plotting residuals 373 against fitted values. For each analysis, we used a model selection process to identify the 374 predictors that best explained variation in the response variable. Model selection was based 375 on minimization of the corrected Akaike's information criterion (AICc) (Burnham and 376 Anderson 2011). Support for an effect of an explanatory variable on the response variable 377 was based on comparison of AICc values between the full model with the effect of interest included vs. excluded, and when $\Delta AICc$ (AICc_{included} – AICc_{excluded}) was less than or equal to 378 379 minus five (Burnham and Anderson 2011). The 95% confidence interval (CI) of the predictor 380 estimates was obtained using the confint function (stats package: R Core Team 2013).

381 The influence of phylogenetic similarity among species was tested in the "best 382 model" obtained during the Imer model selection process (model including only explanatory variables with $\Delta AICc \leq -5$). This was done by running a phylogenetically controlled mixed-383 384 effects model in ASRemI-R (VSN International, Hempstead, U.K.; www.vsn-intl.com) with the 385 same set of predictors as the Imer "best model" for each analysis. The phylogeny was 386 included as a random effect in the form of a correlation matrix of distances from the root of 387 the tree to the most recent common ancestor between two species. The phylogenetic effect 388 was tested by performing a REML likelihood ratio test (comparing the REML likelihood of the

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same ASReml model with and without phylogeny; the log-likelihood ratio test statistic was assessed against a χ^2 distribution with one degree of freedom). The phylogenetic tree used in this comparative study was adapted from a recent species-level molecular phylogenetic assessment (Jetz et al. 2012; Ericson backbone phylogeny) (Figure S1).

393

394 Variation in age at first reproduction

395 To determine how AFR varied within and among species, we noted how often an AFR was 396 the most frequently observed AFR within a species (mode) (Figure S3A) and considered the 397 frequency of a specific AFR age-class across all species (Figure S3B). Then, mean AFR and its 398 standard deviation were calculated for each of the 34 species. We tested the influence of 399 sex, mean lifespan and social lifestyle (family-living and presence of helpers) on variation in 400 mean AFR across the 24 species for which we had data for both sexes (Table S1). We used a 401 weighted linear mixed-effect model with population mean AFR as the unit of analysis, and 402 included species as a random effect. Since AFR cannot exceed the mean lifespan, AFR and 403 mean lifespan should be correlated positively. Therefore, we tested whether the estimated 404 correlation between AFR and mean lifespan differed significantly from the null expectation. 405 To do so, we performed a conservative permutation analysis (following Charmantier et al. 406 2006; Lane et al. 2011). For each mean lifespan, a mean AFR value was randomly selected 407 with replacement from our dataset. During re-sampling we fixed the rule that AFR was 408 smaller than mean lifespan. Data were re-sampled 500 times and analyzed using the same 409 weighted linear mixed-effect model as described above. We estimated the average 410 estimates and 95% Cls over the 500 model outputs and compared them to those observed.

411

412 Fitness consequences of age at first reproduction

413 To determine whether variation in AFR has consequences for LRS, the correlation between 414 AFR and LRS (within-species) as well as its average influence (among-species effect) was 415 investigated using within-subject centering (van de Pol and Wright 2009). The within-species 416 effect was calculated for each sex and species by subtracting the species- and sex-specific 417 mean AFR from each AFR age-class category observed within sex and species (within-species 418 AFR effect; van de Pol and Wright 2009). The among-species effect was determined as the 419 mean AFR within sex and species (between-species AFR effect; van de Pol and Wright 2009). 420 To test for non-linear effects of AFR on LRS within species, a quadratic term of the within-421 species AFR effect was included in the model. The AFR values were centered to reduce collinearity between the within-species AFR effect and the within-species AFR² effect. 422 423 Centering enabled independent interpretation of the linear and the curvature effect 424 (Schielzeth 2010). Due to apparent interspecific variation in the relationship between AFR 425 and LRS, the ideal analytical framework would have been a random intercept and slope 426 model that estimated separate intercepts and slopes for each species. However, our sample size did not provide sufficient power to support such a model (Martin et al. 2011; van de Pol 427 428 2012). Therefore, we ran a standard weighted linear mixed-effect model using the average 429 LRS within AFR age-class categories, with sex and population as units of analysis. Species 430 was included as a random effect in this analysis, along with the natural log of mean body 431 mass as a covariate. We included lifespan in this model as a covariate, since reproductive 432 performance corrected for survival estimates approximates real fitness better (Roff 1992). 433 While the output of the analysis with and without lifespan were similar, lifespan is strongly correlated with the between-species AFR effect. Therefore, we present the analysis without 434 435 lifespan to avoid issues caused by collinearity (Dormann et al. 2013).

To assess whether the most frequently observed AFR within each species was an optimal strategy, the AFR mode within each species was correlated with the AFR that maximized LRS (i.e., Optimal AFR, Table 1). Then, the species-specific Optimal AFR was compared to the age at sexual maturity to identify species with optimal delayed reproduction (i.e., species with Optimal AFR > Age at maturity). Finally, the Optimal AFR was correlated with lifespan to identify if a benefit from delaying the onset of reproduction beyond sexual maturity coincided with long lifespan.

443

444 Among-species variation in the relative timing of optimal age at first reproduction

445 We used a model selection and model averaging approach (Grueber et al. 2011) to 446 determine the factors that explain interspecific variation in deviations of the Optimal AFR 447 from the age of sexual maturity (i.e., Delay Index, Table 1). All life-history, social and 448 ecological factors listed above were included (see DATA COLLECTION), as well as indices 449 reflecting the relationship between LRS and AFR: the Choice Index, and the Before and After 450 Variation Indices (see above, Table 1). In a second analysis, the Lifespan Effect Index was 451 included for the 21 species for which we had detailed data on lifespan mean for each AFR 452 age-class category (Table 1, Figure S2). Due to reduced statistical power of the latter (as on 453 restricted dataset, see above), in the results section we present only the estimates and 95% 454 Cl of the analysis excluding the Lifespan Effect Index. Each of the before mentioned 455 variables, and the biologically relevant interactions (Before Variation Index x After Variation 456 Index, Choice Index x Before Variation Index, Choice Index x After Variation Index, Choice 457 Index x Family-living, Choice Index x Helper presence, Mean lifespan x Family-living, Mean 458 lifespan x Helper presence, Nest predation risk x Family-living, Nest predation risk x Helper 459 presence; Table S8 lists predictions associated with these interactions) were tested against

460 the Delay Index in a weighted linear mixed-effect models with Delay Index for each sex and 461 population as a unit of analysis. Species was added as a random effect. Sex and the natural 462 logarithm of body mass were included as default fixed-effects variables to control for 463 allometry and any differences between sexes. Due to a large number of possible 464 combinations between all predictors, we used the R package MuMIn (Barton 2013) to 465 perform model selection. The candidate model set included models with Δ AICc \leq 5, Δ AICc 466 being the AICc of the focal model minus the AICc of the best model (see Table S9 for analysis 467 excluding Lifespan Effect Index and Table S10 for analysis including Lifespan Effect Index). To 468 estimate the relative importance of a factor, we summed the Akaike's weights of the models 469 in the set of best models including the focal factor, following the method described by Poz 470 Symonds and Moussalli (2011).

471

Results 472

VARIATION IN AGE AT FIRST REPRODUCTION 473

Across species (N = 34), age at first reproduction (AFR) ranged from one to 20 years. In 11 474 475 species, the modal AFR was one year (Figure S3A). In 70% of species, AFR was age 3 or less 476 and only 20% of species had an AFR that was greater than 6 years of age (Figure S3B). 477 Within species, the number of AFR categories ranged from one to 15 (average = 4.8 years; 478 SD = 3.1; N = 34) and the mean AFR and its standard deviation varied among species (Figure 479 1). Removing sex or social variables (i.e. family-living, helper presence) from the model did 480 not influence mean AFR (Table 2). However, mean AFR correlated positively with mean 481 lifespan (parameter estimate for mean lifespan = 0.87, 95% CI (hereafter given in brackets 482 after all estimates): 0.72 to 1.02, Table 2), and this correlation exceeded that expected from 483 the mathematical interdependence of AFR and mean lifespan (estimated by the

permutation test: mean of 500 simulations: 0.63 (0.87 to 0.79), Δ AICc = -22.24). A positive relationship between AFR and mean lifespan was also apparent when comparing the AFR age-class categories within each species (Figure 2). The phylogenetic effect on mean AFR was significant (likelihood ratio test: LRT = 6.99, df = 1, p < 0.01).

488

489 FITNESS CONSEQUENCES OF AGE AT FIRST REPRODUCTION

490 Our within-subject centering approach revealed no among-species effect of AFR on LRS, but a within-species effect of both AFR and AFR² (Figure 3). Within species, there was strong 491 492 directional selection for an early AFR (within-species AFR effect estimate = -0.54 (-0.70 to -0.39), Table S11), as well as stabilizing selection (within-species AFR^2 effect estimate = -0.26 493 (-0.43 to -0.10), Table S11) (Figure 3). The phylogenetic effect on mean LRS for the 494 495 corresponding AFR was not significant (likelihood ratio test: p = 1). Twenty-six out of 34 496 species (76%) had an Optimal AFR delayed beyond the age at maturity, and this delay correlated positively with a longer mean lifespan (slope = 0.28, r_{Spearman} = 0.61, p < 0.005; 497 498 Figure 4). Both the most-observed AFR and mean AFR correlated with the AFR with the 499 highest LRS (Optimal AFR vs. modal AFR: slope = 0.98, $r_{\text{Spearman}} = 0.80$, p < 0.0001; Optimal 500 AFR vs. mean AFR: slope = 0.95, $r_{\text{Spearman}} = 0.84$, p < 0.0001). The latter was true even when 501 only looking at species with a large number of observed AFR age-class categories (Table 502 S12).

503

504 AMONG-SPECIES VARIATION IN THE RELATIVE TIMING OF OPTIMAL AGE AT FIRST 505 REPRODUCTION

506 While the Delay Index was associated with indices that reflect a change in LRS and survival 507 as a function of AFR (i.e. Choice, Before Variation and Lifespan Effect Indices; Table 1, all

508 predictor weights \geq 0.45), it was only marginally related to social (predictor weights < 0.45) 509 or ecological factors (predictor weights \leq 0.30; Tables 3 and 4). A delayed optimal onset of 510 reproduction (i.e. large Delay Index) was found in species with a large range of optimal AFR 511 relative to reproductive lifespan (Choice Index: estimate = 0.44 (0.15 to 0.72), Table 3). 512 Moreover, a large Delay Index was found in species in which early AFR was associated with a 513 decreased LRS (Before Variation Index estimate = 0.30 (0.07 to 0.54), Table 3 and Figure 3) 514 and a reduced reproductive lifespan (Lifespan Effect Index estimate = 0.54 (0.37 to 0.72), 515 Table 4). Finally, larger species showed later optimal onset of reproduction than smaller 516 species (In (body mass) estimate: 0.35 (0.01 to 0.69), Table 3). These results remained 517 quantitatively similar when using indices estimated with the Optimal AFR Range determined 518 under the criterion where AFR categories included in the Optimal AFR Range were AFR(s) 519 adjacent to the Optimal AFR with their 90% Cls overlapping those of the Optimal AFR R. R 520 (Tables S13 to S16).

521

Discussion 522

523 Age at first reproduction (AFR) is a key life-history parameter with consequences for 524 individual reproductive output, and hence its effect on fitness has been studied in a number 525 of intraspecific studies (see references in Table S1). Here we provide a first comparative 526 analysis using a representative amount of averaged within-species information to examine 527 interspecific variation in the relationship between AFR and lifetime reproductive success 528 (LRS). Identifying the species-specific AFR that results in the highest LRS (i.e. Optimal AFR) 529 allowed us to investigate not only within- and among-species variation in the relationship 530 between AFR and LRS, but also differences in the benefits and costs associated with variable 531 timing in the onset of reproduction among species. Our results demonstrated that the most 532 commonly observed AFR within a species corresponds to the species-specific Optimal AFR. 533 Among species, Optimal AFR varied considerably. This study showed that lifespan was a 534 major predictor of the relative timing of the Optimal AFR within the reproductive lifespan 535 and that they correlated positively. Additionally, our analyses revealed that Optimal AFR 536 beyond the age of maturity was associated with a decrease in fitness and survival that arose 537 from starting to reproduce at earlier ages than the Optimal AFR.

538 Age at first reproduction varied considerably both within and among species (Figure 539 1). Some species displayed no variation in AFR (e.g. long-tailed tit Aegithalos caudatus, 540 indigo bunting *Passerina cyanea*, common buzzard *Buteo buteo*), while others exhibited 541 large variation (e.g. mute swan Cygnus olor, wandering albatross Diomedea exulans, 542 eurasian oystercatcher *Haematopus ostralegus*). Most species that expressed variation in 543 AFR experienced negative consequences for LRS from initiating reproduction either too 544 early or too late in life (e.g. the Optimal AFR was at an intermediate point in the 545 reproductive lifespan: between the age of sexual maturity and the oldest AFR observed 546 within a population), while for others the earliest or latest observed AFR resulted in the 547 highest LRS (Figure 3). This suggests simultaneous directional and stabilizing selection. If the 548 pattern observed is a footprint of selection acting at the individual level, this should lead to 549 a decrease in average AFR and a reduction in its evolvability. However, a comparative study 550 directly investigating individual variance would be needed to assess this hipothesis.

551 While there was no overall interspecific relationship between AFR and LRS, a within-552 species relationship between AFR and LRS (Table S11) indicates that evolutionary processes 553 operate at different scales. On the one hand, large-scale evolution acts on all individuals 554 within a population, which might confound the detection of a relationship between AFR and 555 LRS. On the other hand, local-scale evolution acts on individuals, such as on variation in

556 individual quality (Van Noordwijk and De Jong 1986; Kim et al. 2011), food availability 557 (Brommer et al. 1998), territory quality (Krüger 2005), population density (Krüger 2005) or 558 climatic conditions (Gibbs and Grant 1987; Kim et al. 2011), which also might drive the 559 relationship between AFR and LRS. Differences among cohorts in the relationship between 560 AFR and LRS (Brommer et al. 1998; Kim et al. 2011) might additionally explain the absence 561 of a between-species effect of AFR on LRS, but our data did not allow us to take potential 562 differences in individual or cohort quality into account.

563 Among-species variation in mean AFR correlated positively with lifespan (Table 2), 564 supporting the life-history paradigm that the pace of life fundamentally affects reproductive 565 timing (Roff 1992; Stearns 1992; Charlesworth 1994). Furthermore, the species-specific 566 optimal reproductive strategy varied among species, where species with a mean lifespan of 567 up to six years (median mean lifespan: 1.9 years) had an Optimal AFR of one year, providing 568 a quantitative benchmark to differentiate between short- and long-lived bird species. At the 569 other extreme, species with a longer lifespan had a later mean AFR (Table 2) and a later 570 Optimal AFR (Figure 4).

571 When relating the position of the Optimal AFR to the age of sexual maturity of a 572 species, our results revealed that the Optimal AFR was beyond the age of maturity in 26 of 573 34 species. Thus, individuals in these species appear to benefit from delaying their onset of 574 reproduction (e.g. female tawny owl Strix aluco (Millon et al. 2010); female goshawk 575 Accipiter gentilis (Krüger 2005); sexes combined short-tailed shearwater Puffinus tenuirostris 576 (Wooller et al. 1989)). The association of an Optimal AFR beyond the age of sexual maturity 577 with a long mean lifespan suggests that the positive effect of lifespan on mean AFR is not 578 caused by physiological constraints associated with maturity. Indeed, longer-lived species 579 mature later and still adopt an AFR past their age of maturity, and they experienced a larger

LRS as a consequence (Figure 4). Such a benefit from delayed AFR until after the age of sexual maturity was found not only in long-lived species, but also in six out of 11 short-lived species with a mean lifespan of less than three years (Figure 4).

583 When controlling for reproductive lifespan, we found that interspecific variation in 584 deviation of the Optimal AFR from the age at maturity was primarily associated with a 585 change in survival and fitness with AFR (Tables 3 and 4). Moreover, our results confirmed 586 that an early AFR might be favored by a short reproductive lifespan and vice versa (Roff 587 1992; Stearns 1992; Charlesworth 1994) (Table 4 and Figure 2). Species in which an early 588 onset of reproduction was associated with a reduced reproductive lifespan benefited from 589 delaying AFR (Table 4 and Figure S2), which supports the restraint hypothesis (Williams 590 1966; Forslund and Pärt 1995). Moreover, the cost of early reproduction, measured as a 591 decrease in LRS relative to the optimum, correlated positively with the optimal delayed 592 reproductive onset (Table 3). An early reproductive onset might be costly because of 593 differences in individual competitive ability, if this early onset leads to unequal probabilities 594 of acquiring a high-quality territory (Ens et al. 1995; Ekman et al. 2001; Prevot-Julliard et al. 595 2001; Cooper et al. 2009) or to high physiological costs (Hawn et al. 2007). This pattern 596 suggests that different factors affect the evolution of sexual maturity and the onset of 597 reproduction. Interestingly, in species where there was limited change in LRS relative to 598 AFR, postponing the onset of reproduction beyond sexual maturity was chosen over other 599 earlier AFR leading to similar fitness. Therefore, not reproducing as soon as physiologically 600 capable might provide further benefits. Our results provide empirical support for the 601 hypothesis that costs of reproduction shape the onset of reproduction (Lack 1968; Roff 602 1992; Stearns 1992).

603 It has been argued that variation in AFR might be sub-optimal, reflecting constraints 604 on early breeding, such as limited access to high-quality mates or to high-quality breeding 605 sites (Lack 1968; Emlen 1982; Stearns 1989; Koenig et al. 1992). However, our results 606 suggest that the onset of reproduction most likely is an optimal strategy, since the most 607 commonly observed AFR coincides with the Optimal AFR. A number of theories developed 608 to explain the evolution of cooperative breeding depicts the decision of offspring to remain 609 with their parents beyond sexual maturity as a "best of a bad job" strategy that reflects 610 dispersal constraints (Emlen 1982; Koenig et al. 1992; Arnold and Owens 1998; Hatchwell 611 and Komdeur 2000). The lack of a strong correlation between the Delay Index and the 612 different social lifestyles suggests that delayed onset of reproduction might not have 613 evolved due to constraints (Ekman et al. 2004; Ekman 2007), but instead constitutes a 614 beneficial life-history decision, which correlates positively with lifespan (Covas and Griesser 615 2007). Still, the lack of a correlation between social factors and variation in the optimal 616 timing of reproduction could reflect the fact that our data is skewed towards pair-breeding. 617 northern hemisphere species. Including more tropical and southern hemisphere species 618 might alter our results and magnify the role of social factors in our analyses, as the latter 619 two groups are often long-lived (Valcu et al. 2014), stay longer with their parents (Russell 620 2000) and are more likely to breed cooperatively (Jetz and Rubenstein 2011). The current 621 paucity of long-term studies in these regions potentially biases our view of life-history 622 evolution (Martin 2004).

Although we found no significant effect of sex in our study, the relationship between AFR and LRS, and the optimal timing of reproduction, sometimes differed between sexes (Figures 3 and S4). Twelve out of 24 species showed sex-specific differences in the Delay Index; females benefited more from earlier onset than males in seven species, whereas the

627 opposite was true in five species (Figure S4). Intraspecific studies have demonstrated sex 628 differences in the relationship between LRS and AFR (e.g. western gull Larus occidentalis 629 (Pyle et al. 1997); green woodhoopoe Phoeniculus purpureus (Hawn et al. 2007); blue-630 footed booby Sula nebouxii (Kim et al. 2011)), highlighting the need to consider sex-specific 631 variation in life-history traits (McDonald 1993; Santos and Nakagawa 2012). The positive 632 correlation between the relative timing of Optimal AFR and body mass concurs with findings 633 in mammals where AFR is correlated strongly with body mass (larger mammals having later 634 AFR; Estern 1979; Wootton 1987). Nevertheless, we additionally demonstrated that, in 635 birds, larger species benefited more from delaying the onset of reproduction beyond sexual 636 maturity than smaller species. Therefore, body mass seems to be an important factor 637 associated with variation in reproductive strategy. Animals with a large body size invest 638 substantial amounts of resources into growth. Although, in birds, growth after sexual 639 maturity is negligible (Ricklefs 1983), postponing the onset of reproduction might 640 counterbalance the cost endured during the development phase and increase the probability of a high lifetime reproductive output. 641

642 In conclusion, AFR varies both within and among species, and this variation is 643 reflected in LRS. The most frequently observed AFR within a species results in the highest 644 LRS. Where an AFR delayed beyond physiological maturity co-occurred with the highest LRS, 645 this delay was mainly associated with a long lifespan and a decrease in LRS and future 646 survival linked to early reproduction. Our study is the first to provide empirical confirmation 647 of several key predictions of life-history theory across species that lifespan and costs of 648 reproduction shape reproductive timing (Lack 1968; Roff 1992; Stearns 1992; Charlesworth 649 1994). Moreover, the finding that, in long-lived species, postponing the onset of 650 independent reproduction is an optimal strategy has important implications for long-held

perspectives on the evolution of sociality. Hitherto, the decision of young birds to remain with their parents and become helpers has been viewed as a sub-optimal response to the lack of breeding opportunities (Emlen 1982; Koenig et al. 1992; Arnold and Owens 1998). Our results clearly indicate that this decision can be a strategy to mitigate the costs of early reproduction. Overall, our results are consistent with life-history theory and challenge current theories on the evolution of family formation and cooperative breeding.

657

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834 Conflict of Interest

- 835 We have no conflict of interest.
- 836 Author contributions
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- 838 M.V.P., I.G.W., I.S.W., A.W. provided unpublished data on lifetime reproductive success.
- 839 E.M. compiled the data, performed the statistical analysis and wrote the first draft of the

- 840 manuscript. All authors contributed to revisions (especially M.G.) and gave final approval for
- 841 publication. M.G. helped with data compilation and reflection on the manuscript. S.M.D.,
- 842 S.N. and M.G. helped with the statistical methods and estimation of the indices. S.M.D.
- 843 wrote the R script to automate the estimation of two indices.

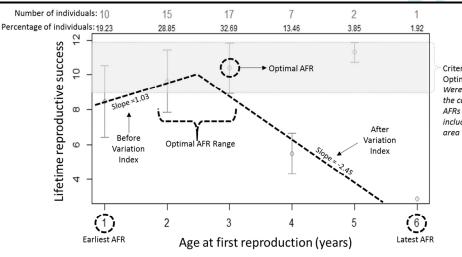
844 Data accessibility

- is 845 The datasets supporting this article have been uploaded as part of the Supporting
- 846 Information.

37

- 847 Table 1. Definitions and descriptions of the parameters and indices estimated for each sex (when
- 848 possible) and each species followed by a graph illustrating the description based on the case of the
- 849 Eurasian sparrowhawk (Accipiter nisus). See also Indices and estimates section in Materials and
- 850 Methods.

Species parameter	Definition	Biological description	Technical description
Optimal AFR	AFR that results in the highest LRS	Reflects the species-average optimum strategy of onset of reproduction	AFR that maximizes mean LRS excluding AFR categories with <10% individuals. Extracted visually
Optimal AFR Range	Range of optimal AFR(s)	Reflects the range of the species- average optimum strategy of onset of reproduction	Number of AFR(s) adjacent to the Optimal AFR with mean LRS values overlapped by the standard error bars of the Optimal AFR. Extracted visually. Range from 1 to 15
Before Variation Index	LRS cost of initiating reproduction before the Optimal AFR Range	Reflects the LRS cost of adopting a reproductive strategy which is earlier than the range of species- average optimum strategy of onset of reproduction	Slope before the Optimal AFR Range (center of the range) between mean LRS and AFR; Average of slopes obtained when all individuals were included, when excluding AFR categories with <5% and <10 % individuals (mean standard error slope = 0.21). A large positive value indicates a strong negative fitness impact of reproducing before the Optimal AFR Range
After Variation Index	LRS cost of initiating reproduction after the Optimal AFR Range	Reflects the LRS cost of adopting a reproductive strategy which is later than the range of species-average optimum strategy of onset of reproduction	Slope after the Optimal AFR Range (center of the range) between mean LRS and AFR; Average of slopes obtained when all individuals were included, when excluding AFR categories with <5% and <10 % individuals (mean standard error slope = 0.18). A large negative value indicates a strong negative fitness impact of reproducing after the Optimal AFR Range
Delay Index	Relative position of the Optimal AFR during the reproductive lifespan	Reflects when – during the average- reproductive lifespan of a species – individuals from a species benefit the most from initiating their reproduction	Varies between 0 and 1. Delay Index 0: the optimal strategy is to start reproduction at physiological maturity; Delay Index 1: the optimal strategy is to delay the onset of reproduction to maximum AFR
Choice Index	Range of optimal AFR(s) relative to the number of AFR observed	Reflects the species-average span of "beneficial choice" in AFR, (i.e. AFRs leading to higher LRS)	Varies between 0 and 1. Choice Index of 0: species has only one optimal AFR; Choice Index of 1: all AFR are optimal
Lifespan Effect Index	Effect of AFR on the mean reproductive lifespan (for each AFR category: see Figure S2)	Reflects the species-specific average effect of the onset of reproduction on survival	Correlation coefficient between mean reproductive lifespan and AFR (Fisher's z transformed) (Koricheva et al. 2013). Positive values suggest a cost of early onset of reproduction, while negative values suggest a cost of late onset



Criterion to estimate the Optimal AFR Range: Were only considered the consecutive adjacent AFRs to the Optimal AFR, included in the shaded area

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Table 2. Effect of sex, mean lifespan of species, family-living and presence of helpers on mean AFR within a species (N = 26 populations, 24 species for which data were available for both sexes). Estimates and 95% confidence intervals (CI) are presented. Δ AIC_c corresponds to the change in AIC_c when the specific parameter was included vs. excluded from the full model.

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50					
57		Standard deviation	Estimate	95% CI	Δ AIC _c
58	Fixed effects:				
59	intercept		0.10	(-0.14, 0.34)	-
60	sex: Female		0.00	na (0.01.0.05)	-2.51
61	sex: Male mean lifespan _{species} *		0.03 0.87	(0.01, 0.05) (0.72, 1.02)	-61.65†
62	family-living: NO		0.00	na	2.58
63	family-living: YES helper presence: NO		-0.12 0.00	(-0.89, 0.64)	
64	helper presence: YES		-0.33	na (-1.16, 0.50)	2.08
65	Random effects:				
	species	0.52		(0.40, 0.70)	
8 9	residuals	0.93		(0.72, 1.26)	
68					

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* factor centered and scaled; na - not applicable; † support for inclusion of the factor

870

871 **Table 3.** Relative importance of predictors included in the full model for the analysis of Delay Index

872 variation excluding Lifespan Effect Index (N = 36 populations, 34 species) and model averaging

estimates (based on 53 models with Δ AICc (AICc focal model – AICc best model) \leq 5, see Table S9).

Predictors	Predict or weight*	Relative importance of predictors†	Model averaging estimates‡,§	95% CI
intercept			0.14	(-0.71, 1.00)
In(body mass)	0.49	1.00	0.35	(0.01, 0.69)
sex	0.49	1.00	Both: 0.00 Female: -0.06 Male: -0.24	na (-0.94, 0.82) (-1.12, 0.65)
Choice Index ¶	0.49	1.00	0.44	(0.15, 0.72)
Before Variation Index ¶	0.48	0.98	0.30	(0.07, 0.54)
family-living	0.40	0.82	NO: 0.00 YES: 0.01	na (-1.45, 1.48)
helper presence	0.40	0.82	NO: 0.00 YES: 0.49	na (-2.12, 3.31)
nest predation risk	0.28	0.56	0.03	(-0.34, 0.43)
Choice Index: helper presence	0.25	0.51	NO: 0.00 YES: -0.67	na (-2.45, -0.18)
mean lifespan	0.25	0.50	0.09	(-0.26, 0.60)
Choice Index: family-living	0.23	0.46	NO: 0.00 YES: 0.57	na (-0.50, 3.00)
mean lifespan: helper presence	0.22	0.44	NO: 0.00 YES: 2.48	na (2.66, 8.49)
mean lifespan: family-living	0.21	0.43	NO: 0.00 YES: -1.91	na (-6.12, -2.72)
nest predation risk: family-living	0.21	0.43	NO: 0.00 YES: 0.91	na (1.23, 2.97)
Before Variation Index: Choice Index	0.17	0.35	0.13	(-0.08, 0.82)
After Variation Index ¶	0.14	0.28	-0.04	(-0.37, 0.05)
nest predation risk: helper presence	0.10	0.21	NO: 0.00 YES: -0.41	na (-3.82, -0.13)
chick development mode	0.05	0.11	Altricial: 0.00 Precocial: -0.02	na (-1.20, 0.74)
latitude	0.03	0.07	-0.01	(-0.41, 0.23)
Before Variation Index: After Variation Index	0.00	0.01	0.00	(-0.08, 0.19)

*: sum of model weights from Table S9 including the focal predictor. na – not applicable;

875 *†*: predictor weight relative to the highest weighted predictor.

876 ‡: model averaging estimates according to full model averaging approach since the best AIC_c model

is not strongly weighted (weight = 0.05) (Symonds and Moussalli 2011).

§: reference levels of categorical variables have an estimate of 0; estimates reflect difference inslope between the reference level and focal level.

880 <u>Note:</u> The relative importance of body mass and sex is due to their inclusion by default in each

881 model to control for allometry and sex differences. All continuous variables are centered and scaled.

882 ¶: predictors reflecting the relationship between LRS and AFR, see Table 1 and the Indices and

883 estimates section of Materials and methods.

Table 4. Relative importance of predictors included in the full model for the analysis of Delay Index

variation including Lifespan Effect Index (N = 22 populations, 21 species) and model averaging

estimates (based on 28 models with \triangle AICc (AICc focal model – AICc best model) \leq 5, see Table S10).

Predictors	Predictor weight*	Relative importance of predictors [†]	Model averaging estimates‡,§	95% CI
intercept			0.42	(-0.34, 1.18)
In(body mass)	0.57	1.00	0.36	(-0.23, 0.95)
sex	0.57	1.00	Both: 0.00 Female: -0.67 Male: -0.35	Na (-1.43, 0.09) (-1.11, 0.42)
Choice Index ¶	0.57	1.00	0.35	(0.17, 0.52)
Lifespan Effect Index ¶	0.57	1.00	0.54	(0.37, 0.72)
After Variation Index ¶	0.38	0.66	-0.20	(-0.56, -0.05)
helper presence	0.26	0.46	NO: 0.00 YES: 0.56	Na (0.03, 2.40)
family-living	0.24	0.42	NO: 0.00 YES: -0.32	Na (-1.59, 0.08)
mean lifespan	0.23	0.41	0.20	(-0.01, 0.98)
chick development mode	0.10	0.17	Altricial: 0.00 Precocial: -0.11	Na (-1.28, 0.04)
nest predation risk	0.10	0.17	0.06	(-0.08, 0.74)
latitude	0.07	0.13	0.02	(-0.32, 0.56)
Choice Index: family-living	0.04	0.07	NO: 0.00 YES: 0.03	Na (-0.14, 0.87)
Choice Index: helper presence	0.04	0.07	NO: 0.00 YES: 0.03	Na (-0.18, 1.07)
Before Variation Index ¶	0.03	0.06	0.01	(-0.16, 0.35)

*: sum of model weights from Table S10 including the focal predictor. na – not applicable;

*: predictor weight relative to the highest weighted predictor.

⁸⁸⁹ ‡: model averaging estimates according to full model averaging approach since the best AIC_c model

is not strongly weighted (weight = 0.10) (Symonds and Moussalli 2011).

§: reference levels of categorical variables have an estimate of 0; estimates reflect difference inslope between the reference level and focal level.

893 <u>Note:</u> The relative importance of body mass and sex is due to their inclusion by default in each 894 model to control for allometry and sex differences. All continuous variables are centered and scaled.

895 ¶: predictors reflecting relationship between LRS or survival and AFR, see Table 1 and the Indices896 and estimates section of Materials and methods.

897 Figure legends

Figure 1. Mean AFR (years) and standard deviation for all 36 populations (34 species) (both sexes combined). Mean AFR ranged from 1 to 12.8 years (mean \pm SD = 3.0 \pm 2.6, N = 36), and standard deviation from 0 to 2.31 (mean \pm SD = 0.80 \pm 0.58, N = 36). A number after the name of a species indicates different populations.

Figure 2. Relationship between AFR (years) and the associated mean lifespan within species and sexes (years, N = 22 populations (21 species) for which detailed data on mean lifespan per AFR category were available). Each point is the mean lifespan of individuals within each AFR category. A number after the name of a species indicates different populations. Regression lines are based on the raw data and were drawn for all cases independent of whether the correlation was significant or not.

Figure 3. Variation in AFR and consequences on fitness - Relationship between standardized LRS and AFR for the 36 populations of the 34 species, separated by sex where possible (a point is the mean LRS (centred and scaled) over all individuals that started to reproduce at a specific AFR). Curves represent quadratic fit of the relationship between standardized LRS and AFR independent of whether the relationship was significant or not.

Figure 4. Species-specific Optimal AFR presented relative to the species age at maturity (left y-axis) with species ordered by mean lifespan (both sexes combined). Mean lifespan values are represented by the grey line and the right y-axis. A number after the name of a species indicates the different populations included in the study.

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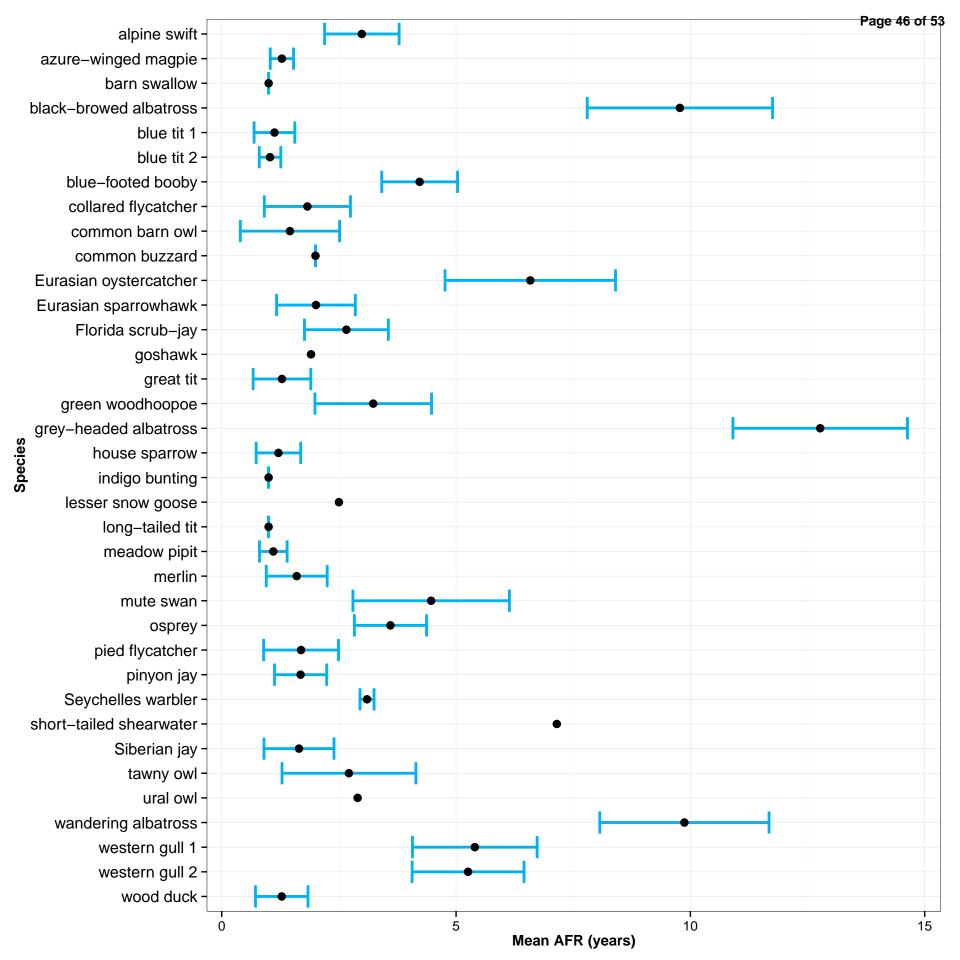
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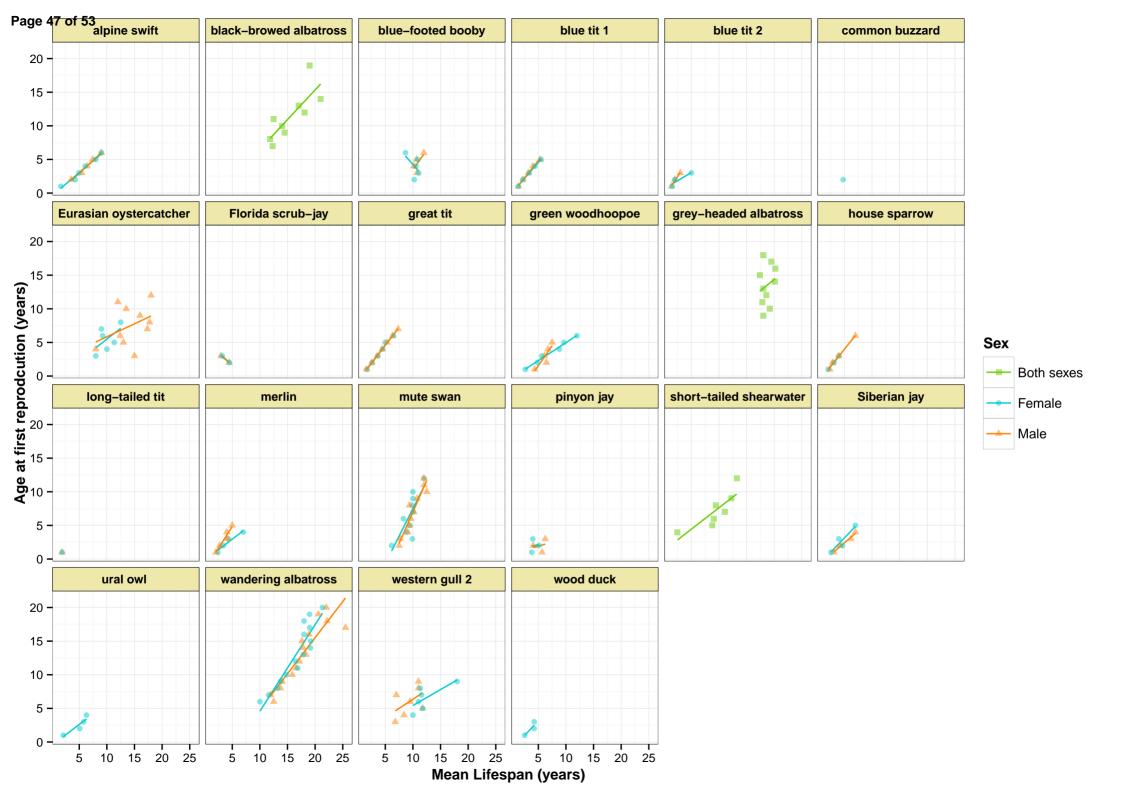
921 Supporting Information

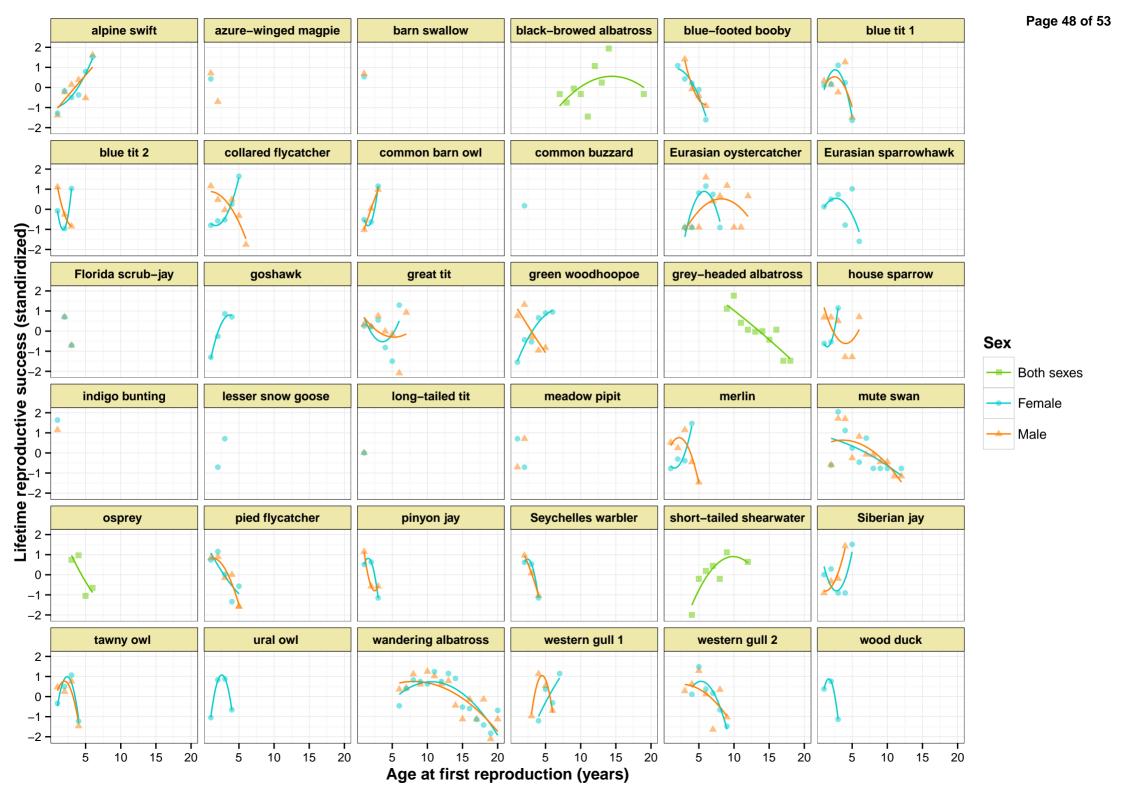
- 922 Additional Supporting Information may be found in the online version of this article at the
- 923 publisher's website:
- 924 **Table S1.** Information on the source and the type of LRS data for each study
- 925 **Table S2.** Model without weighting Variation in AFR analysis
- 926 **Table S3.** Model without weighting Fitness consequence of AFR analysis
- 927 **Table S4.** Model without weighting Delay Index analysis excluding Lifespan Effect Index
- 928 **Table S5.** Model without weighting Delay Index analysis including Lifespan Effect Index
- **Table S6.** Model without weighting Model selection output for the analysis of Delay Index
- 930 variation excluding Lifespan Effect Index
- **Table S7.** Model without weighting Model selection output for the analysis of Delay Index
- 932 variation including Lifespan Effect Index
- 933 **Table S8.** Justification for the interactions used in the analysis of the Delay Index
- **Table S9.** Model selection output for the analysis of Delay Index variation excluding Lifespan
- 935 Effect Index
- 936 Table S10. Model selection output for the analysis of Delay Index variation including
- 937 Lifespan Effect Index
- 938 **Table S11.** Fitness consequence of AFR analysis
- 939 Table S12. Correlation between Optimal AFR vs. modal AFR and mean AFR for different sest
- 940 of species
- 941 Table S13. Model with 90CI Indices Delay Index analysis excluding Lifespan Effect Index
- 942 Table S14. Model with 90CI Indices Delay Index analysis including Lifespan Effect Index

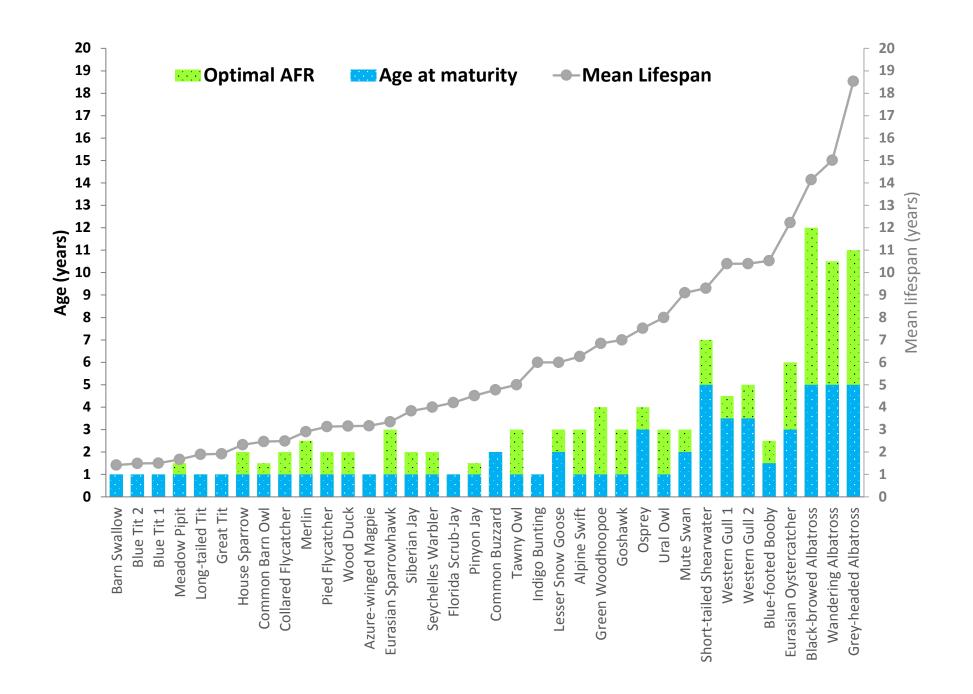
- 943 Table S15. Model with 90Cl Indices – Model selection output for the analysis of Delay Index
- variation excluding Lifespan Effect Index 944
- 945
 Table S16. Model with 90Cl Indices – Model selection output for the analysis of Delay Index
- 946 variation including Lifespan Effect Index
- 947 Figure S1. Phylogenetic tree
- 948 Figure S2. Variation in AFR and consequences on mean reproductive lifespan
- Figure S3. Variation in AFR 949
- 950 Figure S4: Sex differences in the Delay Index

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SUPPORTING INFORMATION

Lifespan and reproductive cost explain interspecific variation in the optimal onset of reproduction

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Table of Contents

Table S1. Information on the source and the type of LRS data for each study.

Species	Scientific name	Location	LRS type	Sample size*	Reference
alpine swift	Apus melba	North-Western Switzerland	fledglings	F: 157; M: 121	Pierre Bize
azure-winged magpie	Cyanopica cyanus	Valdesequera, Spain	fledglings	F: 200; M: 104	Juliana Valencia & Carlos de la Cruz
barn swallow	Hirundo rustica	Kraghede, Denmark	fledglings	F: 1394; M: 1360	Anders Pape Møller
black-browed albatross	Thalassarche melanophris	Bird Island, UK	fledglings	B: 76	Richard Phillips & Andrew G. Wood
blue Tit 1	Cyanistes caeruleus	Oxford, UK	recruits	F: 1177; M: 972	Sandra Bouwhuis & Ben Sheldon
blue Tit 2	Cyanistes caeruleus	Vienna, Austria	recruits	F: 261; M: 211	Bart Kempenaers & Emmi Schlicht
blue-footed booby	Sula nebouxii	Isla Isabela, Mexico	fledglings	F: 222; M: 246	Kim et al. (2011)
collared flycatcher	Ficedula albicollis	Budapest, Hungary	recruits	F: 453; M: 481	Márton Herényi & János Török
common barn owl	Tyto alba	Payerne, Switzerland	fledglings	F: 170; M: 174	Alexandre Roulin
common buzzard	Buteo buteo	Eastern Westphalia, Germany	fledglings	F: 239	Olivier Krüger
Eurasian sparrowhawk	Accipiter nisus	Annandale, Eskdale, Scotland	fledglings	F: 52	McGraw & Caswell (1996)
Eurasian oystercatcher	Haematopus ostralegus	Schiermonnikoog, Netherlands	fledglings	F: 19; M: 33	Martijn Van de Pol
Florida scrub-jay	Aphelocoma coerulescens	Archbold, USA	fledglings	F: 37; M: 43	Fitzpatrick & Woolfenden (1988)
goshawk	Accipiter gentilis	Bissendorf, Spenge, Germany	fledglings	F: 74	Krüger (2005)
great tit	Parus major	Oxford, UK	recruits	F: 4935; M: 4370	Sandra Bouwhuis & Ben Sheldon
green woodhoopoe	Phoeniculus purpureus	Eastern Cape, South Africa	fledglings	F: 59; M: 62	Andrew Radford
grey-headed albatross	Thalassarche chrysostoma	Bird Island, UK	fledglings	B: 74	Richard Phillips & Andrew G. Wood
house sparrow	Passer domesticus	Lundy Island, UK	fledglings	F: 287; M: 265	Terry Burke & colleagues
indigo bunting	Passerina cyanea	Southern Michigan, USA	fledglings	F: 360; M: 357	Payne (1989)
lesser snow goose	Chen caerulescens	La Perouse Bay, Canada	1 st 4 years of life	F: 2616	Viallefont et al. (1995)
long-tailed tit	Aegithalos caudatus	Sheffield, UK	recruits	F: 119; M: 109	Ben Hatchwell
meadow pipit	Anthus pratensis	North-west Germany	fledglings	F: 33; M: 49	Hermann Hötker
merlin	Falco columbarius	Saskatoon, Canada	fledglings	F: 26; M: 68	Richard Espie & Ian G. Warkentin
mute swan	Cygnus olor	Abbotsbury, UK	recruits	F: 252; M: 277	Anne Charmantier, Ben Sheldon & Chris Perrins
osprey	Pandion haliaetus	Michigan, USA	fledglings	B: 40	Postupalsky (1989)
pied flycatcher	Ficedula hypoleuca	Wolfsburg, Germany	fledglings	F: 1411; M: 1135	Sternberg (1989)
pinyon jay	Gymnorhinus cyanocephalus	Flagstaff, USA	yearlings	F: 39; M: 41	John Marzluff
Seychelles warbler	Acrocephalus sechellensis	Cousin Island, Seychelles	fledglings	F: 41; M: 37	Komdeur (1996)
short-tailed shearwater	Puffinus tenuirostris	Fisher Island, Australia	fledglings	B: 186	Wooller et al. (1989)
Siberian jay	Perisoreus infaustus	Arvidsjaur, Sweden	fledglings	F: 44; M: 56	Ekman & Griesser (2016)
tawny owl	Strix aluco	Kielder Forest, UK	fledglings	F: 83; M: 51	Millon et al. (2010)
ural owl	Strix uralensis	Päijät-Häme, Finland	fledglings	F: 57	Brommer et al. (1998)
wandering albatross	Diomedea exulans	Bird Island, UK	fledglings	F: 1819; M: 1519	Richard Phillips & Andrew G. Wood
western gull 1	Larus occidentalis	Farallon Island, USA	fledglings	F: 163; M: 108	Pyle et al. (1997)
western gull 2	Larus occidentalis	Farallon Island, USA	fledglings	F: 66; M: 93	Russell Bradley
wood duck	Aix sponsa	South Carolina, USA	fledglings	F: 90	Oli et al. (2002)

Bold reference indicates unpublished data provided directly by researchers.* number of individuals of F: female, M: male, B: both sexes

Table S2. Model without weighting (see Table 2 for output model with weighting) - Effect of sex, mean lifespan of species, family-living and presence of helpers on mean AFR within a species (N = 26 populations, 24 species for which data were available for both sexes). Estimates and 95% confidence intervals (CI) are presented. Δ AICc corresponds to the change in AICc when the specific parameter was included vs. excluded from the full model.

	Standard deviation	Estimate	95% CI	ΔAIC
Fixed effects:				
intercept		0.10	(-0.14, 0.34)	-
sex: Females		0.00	na	1 00
sex: Male		0.02	(-0.03, 0.07)	1.99
mean lifespan species*		0.82	(0.67, 0.96)	-58.68 ⁻
family-living: NO		0.00	na	
family-living: YES		-0.11	(-0.87 <i>,</i> 0.66)	2.60
helper presence: NO		0.00	na	2.07
helper presence: YES		-0.34	(-1.18, 0.50)	2.07
Random effects:				
species	0.52		(0.40, 0.70)	
residuals	0.09		(0.07, 0.12)	

* factor centered and scaled; na - not applicable; † support for inclusion of the factor

Table S3. **Model without weighting** (see Table S11 for output model with weighting). Results from models testing the within- and among-species effect of AFR on LRS (N = 36 populations, 34 species). Estimates and 95% confidence intervals (CI) are presented. Δ AICc corresponds to the change in AICc when the specific parameter was included vs. excluded from the full model.

	Standard deviation	Estimate	95% CI	Δ AIC _c
Fixed effects:				
intercept		0.00	(-0.36, 0.37)	-
In(body mass)*		-0.01	(-0.17, 0.14)	2.10
sex: Both		0.00	na	
sex: Female		0.12	(-0.28, 0.54)	3.76
sex: Male		0.14	(-0.26, 0.54)	
within-species AFR*		-0.38	(-0.56, -0.20)	-14.97†
within-species AFR ^{2*}		-0.30	(-0.51, -0.10)	-6.36†
between-species AFR		0.03	(-0.16, 0.22)	2.03
Random effects:				
species	0.00		(0.00, 0.13)	
residuals	0.86		(0.79, 0.93)	

* factor centered and scaled; na - not applicable; + support for inclusion of the factor

Table S4. Model without weighting (see Table 3 for output model with weighting). Relative importance of predictors included in the full model for the analysis of Delay Index variation excluding Lifespan Effect Index (N = 36 populations, 34 species) and model averaging estimates (based on 58 models with Δ AICc (AICc focal model – AICc best model) \leq 5, see Table S6).

Predictors	Predict or weight*	Relative importance of predictors†	Model averaging estimates‡,§	95% CI
intercept			0.07	(-0.78, 0.92)
In(body mass)	0.49	1.00	0.36	(0.08, 0.64)
sex	0.49	1.00	Both: 0.00 Female: -0.01 Male: -0.05	na (-0.92, 0.89) (-0.97, 0.87)
Choice Index ¶	0.49	1.00	0.59	(0.34, 0.85)
Before Variation Index ¶	0.48	0.98	0.30	(0.08, 0.52)
Before Variation Index: Choice Index	0.34	0.70	0.25	(0.01, 0.71)
latitude	0.17	0.35	-0.06	(-0.42, 0.06)
After Variation Index ¶	0.12	0.25	-0.03	(-0.31, 0.09)
mean lifespan	0.10	0.20	0.04	(-0.16, 0.54)
family-living	0.08	0.16	NO: 0.00 YES: -0.03	na (-0.65, 0.31)
nest predation risk	0.07	0.14	0.01	(-0.20, 0.31)
helper presence	0.06	0.12	NO: 0.00 YES: -0.00	na (-0.69, 0.66)
chick development mode	0.05	0.11	Altricial: 0.00 Precocial: 0.00	na (-0.75, 0.76)
Choice Index: family-living	0.01	0.03	NO: 0.00 YES: 0.02	na (-0.42, 1.38)
After Variation Index: Choice Index	0.01	0.02	0.00	(-0.15, 0.34)
Before Variation Index: After Variation Index	0.01	0.02	-0.00	(-0.17, 0.11)
mean lifespan: helper presence	0.00	0.01	0.01	(-0.26, 2.12)
Choice Index: helper presence	0.00	0.01	NO: 0.00 YES: -0.01	na (-1.99 <i>,</i> -0.14)

*: sum of model weights from Table S6 including the focal predictor. na – not applicable.

+: predictor weight relative to the highest weighted predictor.

‡: model averaging estimates according to full model averaging approach since the best AIC_c model is not strongly weighted (weight = 0.04) (Symonds and Moussalli 2011).

§: reference levels of categorical variables have an estimate of 0; estimates reflect difference in slope between the reference level and focal level.

<u>Note</u>: The relative importance of body mass and sex is due to their inclusion by default in each model to control for allometry and sex differences. All continuous variables are centered and scaled.

¶: predictors reflecting the relationship between LRS and AFR, see Table 1 and the Indices and estimates section of Materials and methods.

Table S5. Model without weighting (see Table 4 for output model with weighting). Relative importance of predictors included in the full model for the analysis of Delay Index variation including Lifespan Effect Index (N = 22 populations, 21 species) and model averaging estimates (based on 28 models with Δ AICc (AICc focal model – AICc best model) \leq 5, see Table S7).

Predictors	Predictor weight*	Relative importance of predictors†	Model averaging estimates‡,§	95% CI
intercept			0.16	(-0.67 <i>,</i> 0.99)
In(body mass)	0.51	1.00	0.24	(-0.44, 0.92)
			Both: 0.00	na
sex	0.51	1.00	Female: -0.26	(-1.17, 0.65)
			Male: -0.03	(-0.92, 0.87)
Lifespan Effect Index ¶	0.51	1.00	0.54	(0.32, 0.77)
Choice Index ¶	0.49	0.96	0.32	(0.07, 0.60)
mean lifespan	0.30	0.59	0.33	(0.02, 0.10)
After Variation Index ¶	0.25	0.48	-0.13	(-0.51, -0.03)
halmar procence	0.24	0 47	NO: 0.00	na
helper presence		0.47	YES: 0.47	(-0.10, 2.06)
forsily living	0.40	0.25	NO: 0.00	na
family-living	0.18	0.35	YES: -0.26	(-1.57, 0.08)
latitude	0.08	0.15	-0.04	(-0.60, 0.06)
mean lifespan: helper presence	0.05	0.10	0.15	(0.21, 2.82)
Chaine Indow family living	0.05	0.00	NO: 0.00	na
Choice Index: family-living	0.05	0.09	YES: 0.04	(-0.01, 0.95)
Choice Index: helper process	0.04	0.08	NO: 0.00	na
Choice Index: helper presence	0.04	0.08	YES: 0.04	(-0.04, 1.04)
Before Variation Index ¶	0.03	0.05	0.01	(-0.16, 0.34)
shield development mode	0.02	0.05	Altricial: 0.00	na
chick development mode	0.03	0.05	Precocial: -0.02	(-1.00, 0.22)
nest predation risk	0.02	0.03	0.00	(-0.32, 0.34)

*: sum of model weights from Table S7 including the focal predictor. na – not applicable.

+: predictor weight relative to the highest weighted predictor.

‡: model averaging estimates according to full model averaging approach since the best AIC_c model is not strongly weighted (weight = 0.08) (Symonds and Moussalli 2011).

§: reference levels of categorical variables have an estimate of 0; estimates reflect difference in slope between the reference level and focal level.

<u>Note</u>: The relative importance of body mass and sex is due to their inclusion by default in each model to control for allometry and sex differences. All continuous variables are centered and scaled.

¶: predictors reflecting the relationship between LRS or survival and AFR, see Table 1 and the Indices and estimates section of Materials and methods.

	muex	vari	ation exe	Liuuing	<u>s Lite</u>	span	Ellect	muex	lonow	ing on	the nex	t pag	e).												
	Covariate		Ecolog facto		lifes	cial style tors	Life-hi fact	-		ces refle ge in LRS AFR	-				Inte	ractio	ons					Mod	el informa	ation	
(Intercept)	ln(body mass)	sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	After Variation Index : Before Variation Index	After Variation Index : Choice Index	Before Variation Index : Choice Index	Choice Index : helper presence	Choice Index : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Likelihood	AICc	Δ AICc	weight
0.20	0.42	+	-	-	-	-	-	-	0.63	0.30	-	-	-	0.34	-	-	-	-	-	-	9	-68.56	158.58	0.00	0.04
-0.06	0.37	+	-0.17	-	-	-	-	-	0.64	0.34	-	-	-	0.35	-	-	-	-	-	-	10	-67.44	159.19	0.61	0.03
0.14	0.40	+	-	-	-	-	-	-	0.50	0.23	-	-	-	-	-	-	-	-	-	-	8	-70.52	159.75	1.17	0.02
0.05	0.28	+	-	-	-	-	0.20	-	0.65	0.31	-	-	-	0.34	-	-	-	-	-	-	10	-67.80	159.91	1.33	0.02
0.25	0.40	+	-	-	-	-	-	-	0.62	0.31	-0.10	-	-	0.35	-	-	-	-	-	-	10	-68.07	160.45	1.87	0.02
-0.12	0.35	+	-0.17	-	-	-	-	-	0.51	0.27	-	-	-	-	-	-	-	-	-	-	9	-69.52	160.49	1.92	0.02
0.19	0.35	+	-	0.12	-	-	-	-	0.64	0.32	-	-	-	0.41	-	-	-	-	-	-	10	-68.10	160.52	1.94	0.02
-0.03	0.34	+	-0.19	-	-	-	-	-	0.63	0.35	-0.11	-	-	0.36	-	-	-	-	-	-	11	-66.71	160.70	2.12	0.01
0.20	0.42	+	-	-	+	-	-	-	0.62	0.29	-	-	-	0.35	-	-	-	-	-	-	10	-68.29	160.90	2.32	0.01
-0.02	0.25	+	-	-	-	-	0.20	-	0.52	0.25	-	-	-	-	-	-	-	-	-	-	9	-69.77	161.00	2.42	0.01
-0.09	0.37	+	-0.19	-	+	-	-	-	0.63	0.33	-	-	-	0.36	-	-	-	-	-	-	11	-66.94	161.15	2.57	0.01
0.08	0.22	+	-	-	-	-	0.24	-	0.64	0.33	-0.12	-	-	0.35	-	-	-	-	-	-	11	-67.00	161.29	2.71	0.01
0.22	0.41	+	-	-	-	-	-	+	0.63	0.30	-	-	-	0.36	-	-	-	-	-	-	10	-68.54	161.39	2.81	0.01
0.20	0.42	+	-	-	-	+	-	-	0.63	0.30	-	-	-	0.34	-	-	-	-	-	-	10	-68.56	161.43	2.85	0.01
-0.10	0.34	+	-0.21	-	-	+	-	-	0.63	0.33	-	-	-	0.35	-	-	-	-	-	-	11	-67.18	161.64	3.07	0.01
-0.03	0.32	+	-0.19	-	-	-	-	+	0.66	0.34	-	-	-	0.41	-	-	-	-	-	-	11	-67.21	161.70	3.12	0.01
-0.04	0.33	+	-0.15	0.08	-	-	-	-	0.64	0.35	-	-	-	0.39	-	-	-	-	-	-	11	-67.25	161.78	3.20	0.01
0.25	0.29	+	-	0.16	-	-	-	-	0.64	0.35	-0.13	-	-	0.44	-	-	-	-	-	-	11	-67.25	161.78	3.20	0.01
0.17	0.38	+	-	-	-	-	-	-	0.50	0.24	-0.08	-	-	-	-	-	-	-	-	-	9	-70.17	161.81	3.23	0.01
-0.08	0.31	+	-0.13	-	-	-	0.10	-	0.64	0.33	-	-	-	0.35	-	-	-	-	-	-	11	-67.29	161.86	3.28	0.01
-0.01	0.48	+	-	-	-	-	-	-	0.49	-	-	-	-	-	-	-	-	-	-	-	7	-72.98	162.03	3.45	0.01
0.09	0.44	+	-	-	-	-	-	+	0.50	0.24	-	-	-	-	-	-	-	-	-	-	9	-70.30	162.05	3.47	0.01
0.18	0.43	+	-	-	+	-	-	-	0.54	0.28	-	-	-	0.34	-	+	-	-	-	-	11	-67.39	162.06	3.48	0.01
0.13	0.40	+	-	-	+			-	0.50	0.23		-	-	-	-	-	-	-	-	-	9	-70.34	162.15	3.57	0.01

Table S6. Model without weighting (see Table S9 for output model with weighting). Model selection output for the analysis of Delay Index variation excluding Lifespan Effect Index (following on the next page).

	DC		dex vari	ation c		_	Encopa				-		nt page	-]•											
	Covariate		Ecolog facto		Soc lifes fact	tyle	Life-h fact			ces refle ge in LRS AFR				Ir	ntera	ctions						Mod	el informa	ation	
(Intercept)	ln(body mass)	sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	After Variation Index : Before Variation Index	After Variation Index : Choice Index	Before Variation Index : Choice Index	Choice Index : helper presence	Choice Index : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Likihood	AICc	Δ AICc	weight
-0.10	0.32	+	-0.18	-	-	-	-	-	0.50	0.28	-0.10	-	-	-	-	-	-	-	-	-	10	-68.99	162.29	3.71	0.01
0.13	0.39	+	-	0.02	-	-	-	-	0.50	0.24	-	-	-	-	-	-	-	-	-	-	9	-70.50	162.47	3.89	0.01
0.13	0.40	+	-	-	-	+	-	-	0.50	0.23	-	-	-	-	-	-	-	-	-	-	9	-70.51	162.49	3.91	0.01
0.06	0.29	+	-	-	+	-	0.18	-	0.64	0.30	-	-	-	0.35	-	-	-	-	-	-	11	-67.64	162.56	3.98	0.01
-0.15	0.35	+	-0.18	-	+	-	-	-	0.50	0.26	-	-	-	-	-	-	-	-	-	-	10	-69.16	162.64	4.06	0.01
0.01	0.20	+	-	-	-	-	0.24	-	0.52	0.26	-0.11	-	-	-	-	-	-	-	-	-	10	-69.17	162.66	4.08	0.01
-0.06	0.34	+	-0.21	-	+	-	-	-	0.63	0.35	-0.11	-	-	0.37	-	-	-	-	-	-	12	-66.20	162.77	4.19	0.01
0.08	0.28	+	-	0.05	-	-	0.16	-	0.65	0.32	-	-	-	0.37	-	-	-	-	-	-	11	-67.75	162.77	4.20	0.01
-0.08	0.29	+	-0.24	-	-	+	-	-	0.63	0.35	-0.13	-	-	0.37	-	-	-	-	-	-	12	-66.23	162.82	4.24	0.01
0.26	0.40	+	-	-	-	-	-	-	0.62	0.31	-0.12	-	0.10	0.39	-	-	-	-	-	-	11	-67.77	162.83	4.25	0.01
0.06	0.27	+	-	-	-	+	0.20	-	0.65	0.31	-	-	-	0.34	-	-	-	-	-	-	11	-67.79	162.87	4.29	0.00
0.05	0.28	+	-	-	-	-	0.20	+	0.65	0.31	-	-	-	0.34	-	-	-	-	-	-	11	-67.80	162.87	4.29	0.00
0.01	0.27	+	-0.16	0.12	-	-	-	-	0.64	0.38	-0.14	-	-	0.43	-	-	-	-	-	-	12	-66.25	162.88	4.30	0.00
0.24	0.40	+	-	-	+	-	-	-	0.62	0.30	-0.09	-	-	0.36	-	-	-	-	-	-	11	-67.82	162.91	4.33	0.00
0.16	0.46	+	-	-	+	+	-	-	0.62	0.30	-	-	-	0.35	-	-	-	-	-	-	11	-67.85	162.98	4.40	0.00
-0.15	0.33	+	-0.20	-	-	+	-	-	0.51	0.27	-	-	-	-	-	-	-	-	-	-	10	-69.34	162.99	4.41	0.00
-0.14	0.28	+	-0.13	-	-	-	0.11	-	0.52	0.27	-	-	-	-	-	-	-	-	-	-	10	-69.35	163.01	4.43	0.00
0.19	0.36	+	-	0.11	+	-	-	-	0.64	0.32	-	-	-	0.41	-	-	-	-	-	-	11	-67.87	163.01	4.44	0.00
-0.10	0.29	+	-	-	-	-	0.22	+	0.52	0.26	-	-	-	-	-	-	-	-	-	-	10	-69.39	163.08	4.51	0.00
0.11	0.41	+	-	-	+	-	-	-	0.41	0.21	-	-	-	-	-	+	-	-	-	-	10	-69.39	163.09	4.51	0.00
-0.13	0.37	+	-0.16	-	-	-	-	+	0.51	0.27	-	-	-	-	-	-	-	-	-	-	10	-69.46	163.24	4.66	0.00
-0.05	0.25	+	-0.14	-	-	-	0.14	-	0.64	0.35	-0.12	-	-	0.36	-	-	-	-	-	-	12	-66.44	163.25	4.67	0.00
0.25	0.40	+	-	-	-	-	-	-	0.63	0.33	-0.09	-0.03	-	0.35	-	-	-	-	-	-	11	-67.98	163.25	4.67	0.00
-0.01	0.35	+	-0.19	-	-	-	-	-	0.63	0.35	-0.14	-	0.09	0.39	-	-	-	-	-	-	12	-66.44	163.25	4.67	0.00

Table S6 following. Model without weighting (see Table S9 for output model with weighting). Model selection output for the analysis of Delay Index variation excluding Lifespan Effect Index (following on the next page).

Table S6 following. Model without weighting (see Table S9 for output model with weighting). Model selection output for the analysis of Delay Index variation excluding Lifespan Effect Index (following on the next page).

	Covariate			ogical tors	lifes	cial style tors	Life histo facto	ory		ces refle ge in LRS AFR	-			I	ntera	action	IS					Mod	el informa	ation	
(Intercept)	ln(body mass)	sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	After Variation Index : Before Variation Index	After Variation Index : Choice Index	Before Variation Index : Choice Index	Choice Index : helper presence	Choice Index : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Likihood	AICc	A AICc	weight
-0.12	0.36	+	-0.17	-0.02	-	-	-	-	0.51	0.27	-	-	-	-	-	-	-	-	-	-	10	-69.50	163.32	4.74	0.00
0.25	0.39	+	-	-	-	+	-	-	0.62	0.31	-0.10	-	-	0.36	-	-	-	-	-	-	11	-68.06	163.40	4.82	0.00
0.26	0.39	+	-	-	-	-	-	+	0.63	0.31	-0.09	-	-	0.36	-	-	-	-	-	-	11	-68.06	163.41	4.83	0.00
0.00	0.29	+	-0.21	-	-	-	-	+	0.65	0.36	-0.11	-	-	0.41	-	-	-	-	-	-	12	-66.53	163.43	4.85	0.00
0.18	0.36	+	-	0.13	-	-	-	+	0.64	0.33	-	-	-	0.40	-	-	-	-	-	-	11	-68.09	163.46	4.88	0.00
0.40	0.35	+	-	0.12	-	+	-	-	0.64	0.33	-	-	-	0.41	-	-	-	-	-	-	11	-68.10	163.48	4.90	0.00
0.19																					10				
-0.03	0.34	+	-0.19	-	-	-	-	-	0.64	0.37	-0.10	-0.04	-	0.36	-	-	-	-	-	-	12	-66.57	163.50	4.92	0.00
-0.03 0.11	0.34 0.28	+ +	-0.19 -	-	-	- +	- 0.16	-	0.60	0.29	-0.10 -	-0.04 -	-	0.36 0.33	-	-	- +	-	-	-	12	-66.58	163.52	4.94	0.00
-0.03	0.34		-0.19 - -	- - -0.08	- - -	- + -	- 0.16 0.26	- -			-0.10 - -	-0.04 - -	- - -		- -	-	- + -	-	- -	-					

Model set with \triangle AICc \leq 5. N = 36 populations, 34 species.

"+" and "-" indicate the presence or absence of the parameter in the model, respectively. "df" is the degree of freedom. "log Likelihood" is the log likelihood of the model. "AIC_c" represents the Akaike's information criterion corrected for sample size. " Δ AIC_c" is the difference in AIC_c between the focal model and the model with the lowest AIC_c. "weight" represents the relative probability of a model within the full set of models.

	Covariate		Ecolo fact	-	life	ocial style ctors	Life-his facto	-			ng change Il with AF			In	tera	ctior	IS			Мо	del inform	ation	
(Intercept)	ln(body mass)	SeX	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	Lifespan Effect Index	Before Variation Index : Choice	Choice Index : helper presence	Choice Index : family-living	mean lifespan: helper presence	mean lifespan: family-living	nest predation risk : family-living	df	logLikelihood	AICc	A AICc	weight
0.11	-0.07	+	-	-	-	-	0.66	-	0.36	-	-0.27	0.54	-	-	-	-	-	-	10	-30.40	89.61	0.00	0.08
0.31	0.65	+	-	-	+	+	-	-	0.35	-	-	0.57	-	-	-	-	-	-	10	-31.00	90.79	1.19	0.05
0.31	0.60	+	-	-	+	+	-	-	0.20	-	-	0.52	-	-	+	-	-	-	11	-29.15	91.29	1.69	0.04
0.08	0.17	+	-	-	-	-	0.47	-	0.40	-	-	0.56	-	-	-	-	-	-	9	-33.33	91.59	1.99	0.03
0.36	0.56	+	-	-	+	+	-	-	0.22	-	-	0.57	-	+	-	-	-	-	11	-29.38	91.76	2.15	0.03
-0.08	0.43	+	-0.31	-	-	-	-	-	0.40	-	-	0.54	-	-	-	-	-	-	9	-33.56	92.04	2.44	0.02
0.03	-0.03	+	-	-	-	-	0.70	+	0.36	-	-0.33	0.52	-	-	-	-	-	-	11	-29.56	92.12	2.52	0.02
-0.12	-0.03	+	-0.19	-	-	-	0.54	-	0.37	-	-0.27	0.53	-	-	-	-	-	-	11	-29.58	92.16	2.56	0.02
0.18	-0.06	+	-	-	-	+	0.59	-	0.28	-	-0.28	0.51	-	-	-	+	-	-	12	-27.39	92.36	2.75	0.02
0.09	0.01	+	-	-	-	+	0.62	-	0.39	-	-0.27	0.52	-	-	-	-	-	-	11	-29.71	92.41	2.81	0.02
0.45	0.52	+	-	-	-	-	-	-	0.38	-	-	0.56	-	-	-	-	-	-	8	-35.72	92.77	3.16	0.02
0.15	-0.12	+	-	-	-	+	0.64	-	0.25	-	-0.30	0.53	-	+	-	-	-	-	12	-27.68	92.92	3.31	0.02
-0.01	0.36	+	-0.33	-	-	-	-	-	0.37	-	-0.17	0.53	-	-	-	-	-	-	10	-32.30	93.39	3.79	0.01
0.09	-0.09	+	-	-	+	-	0.69	-	0.38	-	-0.29	0.53	-	-	-	-	-	-	11	-30.33	93.65	4.05	0.01
-0.18	0.21	+	-0.21	-	-	-	0.34	-	0.40	-	-	0.54	-	-	-	-	-	-	10	-32.46	93.73	4.12	0.01
0.35	0.42	+	-	-	+	+	0.04	-	-	-	-	0.57	-	-	-	+	-	-	11	-30.36	93.73	4.12	0.01
0.10	-0.08	+	-	0.05	-	-	0.64	-	0.37	-	-0.27	0.56	-	-	-	-	-	-	11	-30.37	93.75	4.14	0.01
0.11	-0.07	+	-	-	-	-	0.66	-	0.36	0.02	-0.27	0.54	-	-	-	-	-	-	11	-30.39	93.77	4.16	0.01
0.34	0.63	+	-	-	+	+	-	-	0.36	0.12	-	0.52	-	-	-	-	-	-	11	-30.43	93.86	4.25	0.01
0.25	-0.19	+	-	-	-	+	0.58	-	-	-	-0.32	0.53	-	-	-	+	-	-	11	-30.46	93.92	4.32	0.01
0.26	0.44	+	-	-	+	+	0.14	-	0.25	-	-	0.55	-	-	-	+	-	-	12	-28.19	93.95	4.34	0.01
0.42	0.46	+	-	-	+	-	-	-	0.14	-	-	0.52	-	-	+	-	-	-	10	-32.59	93.97	4.36	0.01
0.06	0.25	+	-	-	-	+	0.43	-	0.43	-	-	0.53	-	-	-	-	-	-	10	-32.63	94.06	4.45	0.01
0.17	0.48	+	-	-	+	+	0.21	-	0.37	-	-	0.56	-	-	-	-	-	-	11	-30.55	94.09	4.49	0.01

Table S7. Model without weighting (see Table S10 for output model with weighting). Model selection output for the analysis of Delay Index variation including Lifespan Effect Index (following on the next page).

Table S7 following. Model without weighting (see Table S10 for output model with weighting). Model selection output for the analysis of Delay Index variation including Lifespan Effect Index (following on the next page).

	Covariate		Ecolo fact	-	life	ocial style ctors	Life-hi fact	-	:		g change I with AF			In	terac	tion	s			Мо	del inform	ation	
(Intercept)	ln(body mass)	Sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	Lifespan Effect Index	Before Variation Index : Choice	Choice Index : helper presence	Choice Index : family-living	mean lifespan: helper presence	mean lifespan: family-living	nest predation risk : family-living	df	log Likelihood	AICc	A AICc	weight
0.39	0.59	+	-	-	-	+	-	-	0.42	-	-	0.53	-	-	-	-	-	-	9	-34.70	94.33	4.72	0.01
0.25	0.56	+	-	0.13	+	+	-	-	0.37	-	-	0.60	-	-	-	-	-	-	11	-30.73	94.46	4.85	0.01
-0.15	0.37	+	-0.37	-	-	-	-	-	0.42	0.14	-	0.48	-	-	-	-	-	-	10	-32.85	94.50	4.90	0.01
0.37	0.62	+	-	-	+	+	-	-	0.34	-	-0.07	0.56	-	-	-	-	-	-	11	-30.76	94.53	4.92	0.01

Model set with Δ AICc \leq 5. N = 22 populations, 21 species.

"+" and "-" indicate the presence or absence of the parameter in the model, respectively. "df" is the degree of freedom. "log Likelihood" is the log likelihood of the model. " AIC_c " represents the Akaike's information criterion corrected for sample size. " ΔAIC_c " is the difference in AIC_c between the focal model and the model with the lowest AIC_c . "weight" represents the relative probability of a model within the full set of models.

Table S8. Justification for the interactions used in the analysis of the Delay Index.

Interaction	Reason for inclusion in the model
	To test if the relative timing of the Optimal AFR over reproductive lifespan (Delay Index)
Before Variation Index <i>x</i> After Variation Index	was influenced simultaneously by a LRS cost from initiating reproduction both before the
	optimal timing (Before Variation Index) and after the optimal timing (After Variation Index).
	To test if Delay Index was influenced simultaneously by the level of probability to adopt an
	AFR leading to the highest fitness return (i.e. the span of "beneficial AFR" within the
Choice Index x Before Variation Index	observed range of AFR) and a LRS cost from initiating reproduction before Optimal AFR.
	We expect species with a large span of "beneficial AFR" and a low LRS cost of early
	reproduction to benefit from a late AFR.
	To test if Delay Index was influenced simultaneously by the level of probability to adopt an
	AFR leading to the highest fitness return (i.e. the span of "beneficial AFR" within the
Choice Index x After Variation Index	observed range of AFR) and a LRS cost from initiating reproduction after Optimal AFR.
	We expect species with a small span of "beneficial AFR" and a high LRS cost of late
	reproduction to benefit from an early AFR.
Mean lifespan x Family-living	For each of these interactions we tested whether sociality influenced the effect of the focal
	_ predictors on Delay Index based on the idea that living in a kin group (Family living) or
Mean lifespan x Helper presence	breeding cooperatively (Helper presence) might buffer costs associated with the timing of
	— the AFR within the reproductive lifespan.
Nest predation risk x Family-living	For instance, species with a high risk of nest predation need to get experience to successfully
	— defend their nest and have a greater reproductive output. Consequently, they might benefit
Nest predation risk x Helper presence	from a later AFR. However, if the presence of helpers provides anti-predator protection, it
Chaine Index y Family living	<i>— might allow less experienced individuals to still achieve a good reproductive output.</i>
Choice Index x Family-living	Therefore, we expect species with a high risk of nest predation breeding cooperatively to
Choice Index <i>x</i> Helper presence	benefit more from an earlier AFR than species with a high risk of nest predation but
	breeding as a pair without helpers.

The variables included in the interactions are explained in the manuscript as well as in Table 1 for the indices.

	Table	e S9.	Model	selectio	on ou	itput [·]	for the	analy	sis of D	Delay Ir	idex var	iation e	xcludi	ng Life	span	Effe	ect I	nde	x (fc	ollow	ving o	on the ne	ext page).	
	Covariate		Ecolo fact	-	life	cial style tors	Life-hi facto	-		ices refle nge in LR AFR				Inte	ractio	ons						Mo	del inform	ation	
(Intercept)	ln(body mass)	Sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	After Variation Index : Before Variation Index	After Variation Index : Choice Index	Before Variation Index : Choice Index	Choice Index : helper presence	Choice Index : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Likelihood	AICc	Δ AICc	weight
0.14	0.46	+	-	-	+	+	-	-	0.36	0.22	-	-	-	-	+	+	-	-	-	-	12	-60.84	152.05	0.00	0.05
0.06	0.30	+	-	-0.03	+	+	0.19	-	0.36	0.28	-	-	-	-	-	-	+	+	+	+	16	-54.44	152.97	0.92	0.03
0.06	0.33	+	-	-0.03	+	+	0.17	-	0.37	0.28	-	-	-	-	+	-	+	+	-	+	16	-54.74	153.56	1.51	0.02
0.22	0.47	+	-	-	+	+	-	-	0.47	0.29	-	-	-	0.22	+	+	-	-	-	-	13	-60.14	153.86	1.81	0.02
0.23	0.27	+	-	0.28	-	-	-	-	0.67	0.43	-	-	-	0.52	-	-	-	-	-	-	10	-64.84	153.99	1.94	0.02
0.10	0.22	+	-	-0.04	+	+	0.25	-	0.35	0.30	-0.16	-	-	-	-	-	+	+	+	+	17	-53.12	154.16	2.11	0.02
0.19	0.44	+	-	-	+	+	-	-	0.36	0.22	-0.11	-	-	-	+	+	-	-	-	-	13	-60.30	154.19	2.14	0.02
0.33	0.43	+	-	-	-	-	-	-	0.62	0.39	-	-	-	0.42	-	-	-	-	-	-	9	-66.41	154.29	2.24	0.02
0.10	0.25	+	-	-0.04	+	+	0.24	-	0.37	0.29	-0.17	-	-	-	+	-	+	+	-	+	17	-53.28	154.48	2.43	0.01
0.07	0.36	+	-	-0.02	+	+	0.12	-	0.34	0.27	-	-	-	-	-	-	+	+	-	+	15	-57.10	154.64	2.59	0.01
0.10	0.49	+	-	-	+	+	-	+	0.37	0.23	-	-	-	-	+	+	-	-	-	-	13	-60.60	154.79	2.74	0.01
-0.02 0.09	0.40 0.50	+	-0.10	-	+	+	-	-	0.38 0.35	0.24	-	-	-	-	+	+	-	-	-	-	13 11	-60.61 -63.77	154.81 154.82	2.76 2.77	0.01 0.01
0.09	0.30	+ +	-	-	+ +	+ +	- 0.09	-	0.33	- 0.23	_	-	-	-	+ +	+ +	-	-	-	-	13	-60.73	154.82 155.03	2.98	0.01
0.08	0.39	+	_	0.32	-	-	-	_	0.69	0.23	-0.15	_	-	0.57	т -	т -	_	_	_	_	11	-63.89	155.06	3.01	0.01
0.06	0.32	+	-	-0.03	+	+	0.17	_	0.37	0.29	-	-	-	-	-	+	+	+	_	+	16	-55.49	155.06	3.01	0.01
0.13	0.45	+	-	0.02	+	+	-	_	0.37	0.22	-	-	-	-	+	+	-	-	-	-	13	-60.83	155.25	3.20	0.01
0.15	0.29	+	-	0.03	+	+	0.14	-	0.46	0.35	-	-	-	0.23	_	_	+	+	+	+	17	-53.73	155.38	3.33	0.01
0.05	0.23	+	-	0.03	+	+	0.22	+	0.36	0.33	-0.22	-	-	-	-	-	+	+	+	+	18	-51.82	155.55	3.50	0.01
0.05	0.28	+	-	-0.03	+	+	0.21	-	0.38	0.29	-	-	-	-	-	+	+	+	+	+	17	-53.84	155.60	3.55	0.01
0.30	0.44	+	-	-	+	+	-	-	0.47	0.31	-0.13	-	-	0.25	+	+	-	-	-	-	14	-59.38	155.69	3.64	0.01
0.10	0.23	+	-	-0.04	+	+	0.25	-	0.37	0.30	-0.17	-	-	-	-	+	+	+	-	+	17	-54.01	155.92	3.87	0.01
0.22	0.20	+	-	0.04	+	+	0.21	-	0.48	0.38	-0.18	-	-	0.28	-	-	+	+	+	+	18	-52.01	155.93	3.88	0.01
0.06	0.30	+	-	-0.03	+	+	0.19	-	0.37	0.28	-	-	-	-	+	-	+	+	+	+	17	-54.04	155.99	3.94	0.01
0.11	0.29	+	-	-0.03	+	+	0.19	-	0.33	0.28	-0.15	-	-	-	-	-	+	+	-	+	16	-55.97	156.03	3.98	0.01

	Covariate			ogical tors	life	cial style ctors	Life-his facto			ces refle ge in LR: AFR	-			Inte	ractio	ons						Mod	lel inform	ation	
(Intercept)	ln(body mass)	SeX	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	After Variation Index : Before Variation Index	After Variation Index : Choice Index	Before Variation Index : Choice Index	Choice Index : helper presence	Choice Index : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Likelihood	AICc	Δ AICc	weight
0.02	0.32	+	-	0.01	+	+	0.16	+	0.36	0.30	-	_	-	-	-	-	+	+	+	+	17	-54.06	156.03	3.98	0.01
0.44	0.35	+	-	-	-	-	-	+	0.65	0.40	-	-	-	0.48	-	-	-	-	-	-	10	-65.86	156.04	3.99	0.01
0.15	0.32	+	-	0.03	+	+	0.12	-	0.47	0.35	-	-	-	0.22	+	-	+	+	-	+	17	-54.07	156.05	4.00	0.01
0.10	0.39	+	-	-	+	+	0.07	-	0.37	0.22	-	-	-	-	+	+	+	-	-	-	14	-59.66	156.25	4.20	0.01
0.39	0.42	+	-	-	-	-	-	-	0.63	0.40	-0.11	-	-	0.44	-	-	-	-	-	-	10	-65.98	156.27	4.22	0.01
0.22	0.22	+	-	0.03	+	+	0.19	-	0.49	0.38	-0.19	-	-	0.28	+	-	+	+	-	+	18	-52.21	156.33	4.28	0.01
0.06	0.26	+	-	0.02	+	+	0.21	+	0.37	0.33	-0.22	-	-	-	+	-	+	+	-	+	18	-52.22	156.34	4.29	0.01
0.10	0.39	+	-0.13	-	-	-	-	-	0.63	0.41	-	-	-	0.42	-	-	-	-	-	-	10	-66.04	156.38	4.33	0.01
0.18	0.35	+	-	0.05	+	+	0.07	-	0.46	0.35	-	-	-	0.27	-	-	+	+	-	+	16	-56.16	156.41	4.36	0.01
0.16	0.48	+	-	-	+	+	-	+	0.36	0.24	-0.14	-	-	-	+	+	-	-	-	-	14	-59.74	156.42	4.37	0.01
0.25	0.20	+	-	0.05	+	+	0.20	-	0.53	0.42	-0.20	-	-	0.35	-	+	+	+	-	+	18	-52.32	156.54	4.49	0.01
-0.02	0.34	+	-0.15	-	+	+	-	-	0.38	0.25	-0.14	-	-	-	+	+	-	-	-	-	14	-59.81	156.57	4.52	0.01
0.09	0.20	+	-	-0.04	+	+	0.28	-	0.37	0.31	-0.17	-	-	-	-	+	+	+	+	+	18	-52.33	156.58	4.53	0.00
0.05	0.40	+	-0.11	-	+	+	-	-	0.48	0.31	-	-	-	0.22	+	+	-	-	-	-	14	-59.86	156.66	4.61	0.00
0.18	0.30	+	-	0.05	+	+	0.12	-	0.50	0.38	-	-	-	0.29	-	+	+	+	-	+	17	-54.38	156.67	4.62	0.00
0.30	0.24	+	-	0.25	-	-	-	+	0.68	0.43	-	-	-	0.55	-	-	-	-	-	-	11	-64.70	156.67	4.62	0.00
0.18	0.42	+	-	-	-	-	-	-	0.43	0.25	-	-	-	-	-	-	-	-	-	-	8	-68.98	156.67	4.62	0.00
0.20	0.41	+	-	0.10	+	+	-	-	0.49	0.32	-	-	-	0.27	+	+	-	-	-	-	14	-59.91	156.76	4.71	0.00
0.09	0.30	+	0.03	-0.03	+	+	0.20	-	0.36	0.28	-	-	-	-	-	-	+	+	+	+	17	-54.43	156.76	4.71	0.00
0.30	0.31	+	-	0.32	-	-	-0.09	-	0.66	0.44	-	-	-	0.53	-	-	-	-	-	-	11	-64.74	156.77	4.72	0.00
0.13	0.26	+	-0.06	0.26	-	-	-	-	0.67	0.44	-	-	-	0.52	-	-	-	-	-	-	11	-64.75	156.77	4.72	0.00
0.19	0.44	+	-	-	+	+	-	-	0.36	0.17	-0.14	0.06	-	-	+	+	-	-	-	-	14	-59.94	156.82	4.77	0.00

Table S9 following. Model selection output for the analysis of Delay Index variation excluding Lifespan Effect Index.

	Covariate			ogical ctors	life	cial style tors	Life-his facto			ces refle ge in LRS AFR				Inte	ractio	ns						Mod	el informa	ation	
(Intercept)	ln(body mass)	sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	After Variation Index : Before Variation Index	After Variation Index : Choice Index	Before Variation Index : Choice Index	Choice Index : helper presence	Choice Index : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Likelihood	AICc	Δ ΑΙСС	weight
0.04	0.35	+	-	0.00	+	+	0.15	+	0.37	0.29	-	-	-	-	+	-	+	+	-	+	17	-54.49	156.89	4.84	0.00
0.16	0.38	+	-	-	+	+	0.10	-	0.48	0.30	-	-	-	0.22	+	+	-	-	-	-	14	-59.98	156.90	4.85	0.00
0.25	0.37	+	-	-	-	-	0.09	-	0.63	0.39	-	-	-	0.42	-	-	-	-	-	-	10	-66.30	156.91	4.86	0.00
0.23	0.27	+	-	0.28	-	+	-	-	0.67	0.44	-	-	-	0.52	-	-	-	-	-	-	11	-64.83	156.93	4.88	0.00
0.24	0.27	+	-	0.28	+	-	-	-	0.67	0.44	-	-	-	0.52	-	-	-	-	-	-	11	-64.83	156.95	4.90	0.00
0.11	0.32	+	-	-	+	+	0.14	-	0.37	0.23	-0.13	-	-	-	+	+	-	-	-	-	14	-60.03	157.00	4.95	0.00

Table S9 following. Model selection output for the analysis of Delay Index variation excluding Lifespan Effect Index.

Model set with \triangle AICc \leq 5. N = 36 populations, 34 species.

"+" and "-" indicate the presence or absence of the parameter in the model, respectively. "df" is the degree of freedom. "log Likelihood" is the log likelihood of the model. "AIC_c" represents the Akaike's information criterion corrected for sample size. " Δ AIC_c" is the difference in AIC_c between the focal model and the model with the lowest AIC_c. "weight" represents the relative probability of a model within the full set of models.

Table S10. Model selection output for the analysis of Delay Index variation including Lifespan Effect Index.

	Covariate			ogical tors	life	ocial style ctors	Life-his facto	•			g change I with AF			In	itera	ctior	IS			Мос	lel inform	ation	
(Intercept)	ln(body mass)	sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	Lifespan Effect Index	Before Variation Index : Choice	Choice Index : helper presence	Choice Index : family-living	mean lifespan: helper presence	mean lifespan: family-living	nest predation risk : family-living	df	log Likelihood	AICc	Δ AICc	weight
0.36	0.05	+	-	-	-	-	0.53	-	0.36	-	-0.33	0.53	-	-	-	-	-	-	10	-24.97	78.73	0.00	0.10
0.42	0.61	+	-	-	+	+	-	-	0.35	-	-	0.53	-	-	-	-	-	-	10	-25.55	79.90	1.17	0.06
0.35	0.30	+	-	0.39	-	-	-	+	0.38	-	-0.36	0.61	-	-	-	-	-	-	11	-23.91	80.82	2.09	0.04
0.51	0.58	+	-	-	+	+	-	-	0.33	-	-0.19	0.52	-	-	-	-	-	-	11	-23.91	80.82	2.09	0.04
0.35	0.08	+	-	-	-	+	0.53	-	0.38	-	-0.32	0.51	-	-	-	-	-	-	11	-24.30	81.61	2.87	0.02
0.68	0.78	+	0.21	-	+	+	-	-	0.31	-	-	0.54	-	-	-	-	-	-	11	-24.39	81.78	3.05	0.02
0.41	0.16	+	-	-	-	-	0.43	+	0.35	-	-0.36	0.52	-	-	-	-	-	-	11	-24.42	81.85	3.11	0.02
0.28	0.00	+	-	0.18	-	-	0.46	-	0.38	-	-0.33	0.57	-	-	-	-	-	-	11	-24.48	81.97	3.23	0.02
0.60	0.55	+	-	-	-	-	-	+	0.32	-	-0.33	0.54	-	-	-	-	-	-	10	-26.64	82.09	3.36	0.02
0.41	0.59	+	-	-	+	+	-	-	0.31	-	-	0.51	-	-	+	-	-	-	11	-24.69	82.38	3.65	0.02
0.42	0.59	+	-	-	+	+	-	-	0.31	-	-	0.52	-	+	-	-	-	-	11	-24.70	82.41	3.68	0.02
0.29	0.24	+	-	-	-	-	0.37	-	0.38	-	-	0.56	-	-	-	-	-	-	9	-28.90	82.72	3.98	0.01
0.33	-0.01	+	-	-	+	-	0.60	-	0.37	-	-0.35	0.52	-	-	-	-	-	-	11	-24.88	82.76	4.03	0.01
0.33	0.37	+	-0.21	-	-	-	-	-	0.35	-	-0.28	0.55	-	-	-	-	-	-	10	-27.02	82.83	4.10	0.01
0.35	0.04	+	-	-	-	-	0.54	-	0.36	0.02	-0.33	0.52	-	-	-	-	-	-	11	-24.95	82.90	4.17	0.01
0.37	0.04	+	0.02	-	-	-	0.56	-	0.36	-	-0.33	0.53	-	-	-	-	-	-	11	-24.96	82.91	4.18	0.01
0.75	0.80	+	0.28	-	+	+	-	-	0.24	-	-	0.51	-	-	+	-	-	-	12	-22.69	82.94	4.20	0.01
0.41	0.61	+	-	-	+	+	-	-	0.35	0.11	-	0.52	-	-	-	-	-	-	11	-25.04	83.07	4.34	0.01
0.48	0.55	+	-	-	-	-	-	-	0.35	-	-	0.59	-	-	-	-	-	-	8	-30.90	83.14	4.41	0.01
0.33	0.33	+	-	0.40	-	+	-	+	0.41	-	-0.36	0.58	-	-	-	-	-	-	12	-22.80	83.17	4.44	0.01

Table S10 following. Model selection output for the analysis of Delay Index variation including Lifespan Effect Index.

	Covariate			ogical tors	life	ocial style ctors	Life-hi facto	•			ng change Il with AF			In	itera	ction	s			Мо	del inform	ation	
(Intercept)	ln(body mass)	sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index	Before Variation Index	After Variation Index	Lifespan Effect Index	Before Variation Index : Choice	Choice Index : helper presence	Choice Index : family-living	mean lifespan: helper presence	mean lifespan: family-living	nest predation risk : family-living	df	log Likelihood	AICc	ΔAICc	weight
0.50	0.56	+	-	-	+	+	-	-	0.29	-	-0.20	0.50	-	-	+	-	-	-	12	-22.84	83.24	4.51	0.01
0.29	0.26	+	-	0.45	-	-	-	+	0.39	0.17	-0.38	0.59	-	-	-	-	-	-	12	-22.84	83.25	4.52	0.01
0.75	0.78	+	0.27	-	+	+	-	-	0.25	-	-	0.53	-	+	-	-	-	-	12	-22.85	83.26	4.53	0.01
0.51	0.55	+	-	-	+	+	-	-	0.29	-	-0.19	0.52	-	+	-	-	-	-	12	-22.89	83.35	4.62	0.01
0.35	0.27	+	-	0.37	-	-	-	-	0.38	-	-0.24	0.65	-	-	-	-	-	-	10	-27.28	83.37	4.64	0.01
0.52	0.80	+	-	-	+	+	-0.21	-	0.33	-	-	0.54	-	-	-	-	-	-	11	-25.27	83.53	4.80	0.01
0.34	0.53	+	-	0.13	+	+	-	-	0.37	-	-	0.56	-	-	-	-	-	-	11	-25.31	83.61	4.88	0.01
0.58	0.52	+	-	-	-	-	-	-	0.33	-	-0.21	0.58	-	-	-	-	-	-	9	-29.39	83.69	4.96	0.01

Model set with Δ AIC_c \leq 5. N = 22 populations, 21 species.

"+" and "-" indicate the presence or absence of the parameter in the model, respectively. "df" is the degree of freedom. "log Likelihood" is the log likelihood of the model. " AIC_c " represents the Akaike's information criterion corrected for sample size. " ΔAIC_c " is the difference in AIC_c between the focal model and the model with the lowest AIC_c . "weight" represents the relative probability of a model within the full set of models.

Table S11. Results from models testing the within- and among-species effect of AFR on LRS (N = 36 populations, 34 species). Estimates and 95% confidence intervals (CI) are presented. Δ AICc corresponds to the change in AICc when the specific parameter was included vs. excluded from the full model.

	Standard deviation	Estimate	95% CI	Δ AIC _c
Fixed effects:				
intercept		-0.12	(-0.87, 0.62)	-
In(body mass)*		-0.11	(-0.29, 0.08)	0.91
sex: Both		0.00	na	
sex: Female		0.38	(-0.37, 1.14)	3.09
sex: Male		0.41	(-0.34, 1.16)	
within-species AFR*		-0.54	(-0.70, -0.39)	-43.83
within-species AFR ^{2*}		-0.26	(-0.43, -0.10)	-7.45†
between-species AFR		0.08	(-0.13, 0.29)	1.57
Random effects:				
species	0.00		(0.00, 0.16)	
residuals	19.64		(18.14, 21.35)	

* factor centered and scaled; na - not applicable; † support for inclusion of the factor

Table S12. Correlation between Optimal AFR vs. modal AFR and Optimal AFR vs. mean AFR.

	Mean	AFR vs. Optimal	AFR	AFR mode vs. Optimal AFR								
cases	Correlation coefficient	Р	Slope	Correlation coefficient	Ρ	Slope						
all	0.84	< 0.0001	0.95	0.80	< 0.0001	0.98						
(N=62)	(Spearman)		0.00	(Spearman)		0.50						
with AFR range > 4 (N=29)	0.85 (Spearman)	< 0.0001	0.99	0.87 (Spearman)	< 0.0001	0.82						
with AFR range > 6 (N=12)	0.96 (Pearson)	< 0.0001	1.13	0.92 (Pearson)	< 0.0001	1.06						

Table S13. Model with 90Cl indices (see Table 3 for comparison). Relative importance of predictors included in the full model for the analysis of Delay Index variation excluding Lifespan Effect Index (N = 36 populations, 34 species) and model averaging estimates (based on 51 models with Δ AICc (AICc focal model – AICc best model) \leq 5, see Table S15).

Predictors	Predict or weight*	Relative importance of predictors ⁺	Model averaging estimates‡,§	95% CI
intercept		-	0.42	(-0.60, 1.44)
In(body mass)	0.51	1.00	0.21	(-0.17, 0.59)
			Both: 0.00	na
sex	0.51	1.00	Female: -0.30	(-1.40 <i>,</i> 0.79)
			Male: -0.54	(-1.64, 0.56)
Choice Index 90Cl ¶	0.51	1.00	0.53	(0.29, 0.77)
Before Variation Index 90CI ¶	0.51	1.00	0.36	(-0.02, 0.74)
Before Variation Index 90CI: Choice Index 90CI	0.51	1.00	0.86	(0.35, 1.37)
nest predation risk	0.41	0.80	0.35	(0.03, 0.84)
After Variation Index 90CI ¶	0.35	0.69	-0.17	(-0.49, 0.01)
Before Variation Index 90CI: After Variation Index 90CI	0.17	0.33	-0.05	(-0.32, 0.01)
family-living	0.13	0.26	NO: 0.00	na
ianny-nving	0.13	0.20	YES: 0.15	(-0.95, 2.15)
chick development mode	0.12	0.24	Altricial: 0.00	na
	0.12	0.24	Precocial: 0.17	(-0.28, 1.69)
mean lifespan	0.10	0.20	-0.04	(-0.77, 0.35)
helper presence	0.10	0.19	NO: 0.00	na
	0.10	0.19	YES: -0.12	(-2.91, 1.63)
nest predation risk: family-living	0.05	0.09	NO: 0.00	na
	0.05	0.09	YES: 0.09	(-0.34, 2.21)
Choice Index: helper presence	0.04	0.08	NO: 0.00	na
choice index. helper presence	0.04	0.08	YES: -0.12	(-2.55, -0.30)
latitude	0.04	0.08	0.00	(-0.30, 0.42)
Choice Index: family-living	0.04	0.07	NO: 0.00	na
choice index. fairing-ining	0.04	0.07	YES: 0.11	(0.62, 2.44)
After Variation Index 90CI: Choice Index 90CI	0.02	0.04	-0.00	(-0.53, 0.48)
mean lifespan: helper presence	0.02	0.03	NO: 0.00	na
	0.02	0.05	YES: 0.13	(2.18, 7.04)
mean lifespan: family-living	0.02	0.03	NO: 0.00	na
	0.02	0.05	YES: -0.10	(-5.17, -1.96)
nest predation risk: helper presence	0.01	0.01	NO: 0.00	na
חנשנ פרכתמנוטוו וושא. חפופיר פרפשרונכ	0.01	0.01	YES: -0.01	(-3.41, 0.27)

*: sum of model weights from Table S15 including the focal predictor. na – not applicable.

+: predictor weight relative to the highest weighted predictor.

‡: model averaging estimates according to full model averaging approach since the best AIC_c model is not strongly weighted (weight = 0.05) (Symonds and Moussalli 2011).

§: reference levels of categorical variables have an estimate of 0; estimates reflect difference in slope between the reference level and focal level.

<u>Note:</u> The relative importance of body mass and sex is due to their inclusion by default in each model to control for allometry and sex differences. All continuous variables are centered and scaled.

¶: predictors reflecting the relationship between LRS and AFR, see Table 1 and the Indices and estimates section of Materials and methods.

Table S14. Model with 90CI indices (see Table 4 for comparison). Relative importance of predictors included in the full model for the analysis of Delay Index variation including Lifespan Effect Index (N = 22 populations, 21 species) and model averaging estimates (based on 45 models with Δ AICc (AICc focal model – AICc best model) \leq 5, see Table S16).

Predictors	Predictor weight*	Relative importance of predictors†	Model averaging estimates‡,§	95% CI
intercept			0.64	(-0.24, 1.52)
In(body mass)	0.61	1.00	0.47	(-0.02, 0.96)
			Both: 0.00	na
sex	0.61	1.00	Female: -0.82	(-1.51, 0.34)
			Male: -0.59	(-1.52, 0.34)
Lifespan Effect Index ¶	0.61	1.00	0.47	(0.25, 0.69)
Choice Index 90CI ¶	0.58	0.96	0.34	(0.11, 0.61)
Before Variation Index 90Cl ¶	0.32	0.52	0.12	(-0.09, 0.56)
Before Variation Index 90CI: Choice Index 90CI	0.31	0.51	0.25	(0.18, 0.83)
After Variation Index 90Cl ¶	0.29	0.47	-0.12	(-0.51, -0.01)
family living	0.16	0.27	NO: 0.00	na
family-living	0.10	0.27	YES: -0.25	(-2.30, 0.45)
nest predation risk	0.15	0.25	0.10	(-0.01, 0.82)
helper processo	0.15	0.25	NO: 0.00	na
helper presence	0.15	0.25	YES: 0.42	(-1.19, 4.58)
	0.00	0.45	Altricial: 0.00	na
chick development mode	0.09	0.15	Precocial: -0.08	(-1.27, 0.18)
mean lifespan	0.08	0.14	-0.02	(-0.94, 0.71)
latitude	0.08	0.14	0.03	(-0.33, 0.70)
	0.04	0.00	NO: 0.00	na
Mean lifespan: helper presence	0.04	0.06	YES: 0.20	(-0.44, 7.34)
Chains Indow halman processo	0.01	0.03	NO: 0.00	na
Choice Index: helper presence	0.01	0.02	YES: -0.03	(-2.45, -0.28)
Choice Indow family living	0.01	0.02	NO: 0.00	na
Choice Index: family-living	0.01	0.02	YES: 0.01	(-0.11, 0.80)

*: sum of model weights from Table S16 including the focal predictor. na – not applicable.

+: predictor weight relative to the highest weighted predictor.

‡: model averaging estimates according to full model averaging approach since the best AIC_c model is not strongly weighted (weight = 0.06) (Symonds and Moussalli 2011).

§: reference levels of categorical variables have an estimate of 0; estimates reflect difference in slope between the reference level and focal level.

<u>Note:</u> The relative importance of body mass and sex is due to their inclusion by default in each model to control for allometry and sex differences. All continuous variables are centered and scaled.

¶: predictors reflecting the relationship between LRS or survival and AFR, see Table 1 and the Indices and estimates section of Materials and methods.

	excluding Lifespan Effect index (following on the next page).																								
	Covariate		Ecolog facto	-	life	cial style ctors	Life-his facto		:	ces refle ge in LR AFR	-			Inte	eracti	ons			Model information						
(Intercept)	ln(body mass)	SeX	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index 90Cl	Before Variation Index 90Cl	After Variation Index 90Cl	After Variation Index 90Cl : Before Variation Index 90Cl	After Variation Index 90Cl : Choice Index 90Cl	Before Variation Index 90Cl : Choice Index 90Cl	Choice Index 90Cl : helper presence	Choice Index 90CI : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Likelihood	AICc	Δ AICc	weight
0.27	0.16	+	-	0.44	-	-	-	-	0.52	0.31	-0.25	-	-	0.81	-	-	-	-	-	-	11	-67.68	162.63	0.00	0.04
0.32	0.13	+	-	0.49	-	-	-	-	0.59	0.56	-0.25	-0.14	-	1.03	-	-	-	-	-	-	12	-66.22	162.80	0.17	0.04
0.22	0.25	+	-	0.35	-	-	-	-	0.47	0.23	-	-	-	0.71	-	-	-	-	-	-	10	-69.72	163.76	1.12	0.02
0.57	0.29	+	-	0.62	-	-	-0.33	-	0.61	0.60	-0.23	-0.17	-	1.11	-	-	-	-	-	-	13	-65.15	163.89	1.26	0.02
0.63	0.29	+	-	-	-	-	-	+	0.49	0.19	-	-	-	0.76	-	-	-	-	-	-	10	-69.83	163.98	1.34	0.02
0.47	0.18	+	-	0.27	-	-	-	+	0.51	0.21	-	-	-	0.80	-	-	-	-	-	-	11	-68.66	164.61	1.97	0.02
0.44	0.27	+	-	0.53	-	-	-0.23	-	0.52	0.30	-0.24	-	-	0.83	-	-	-	-	-	-	12	-67.19	164.74	2.11	0.02
0.50	0.09	+	-	0.36	+	-	-	-	0.55	0.29	-0.27	-	-	0.88	-	-	-	-	-	+	13	-65.61	164.81	2.17	0.01
0.42	0.12	+	-	0.37	-	-	-	+	0.53	0.29	-0.22	-	-	0.85	-	-	-	-	-	-	12	-67.26	164.88	2.24	0.01
0.48 0.35	0.09 0.45	+	-	0.41	-	-	-	+	0.61 0.43	0.54 0.22	-0.21	-0.15	-	1.09 0.62	-	-	-	-	-	-	13	-65.68 -71.77	164.94 165.01	2.30 2.37	0.01
0.35	0.43	+ +	- 0.11	- 0.48	-	-	-	-	0.43	0.22	- -0.25	-	-	0.82	-	-	-	-	-	-	9 12	-67.46	165.01	2.57	0.01 0.01
0.48	0.17	+	-	0.48 0.46	-	-	- -0.28	-	0.55	0.51	-0.25	-	-	0.85	-	-	-	-	-	-	12	-67.40 -69.01	165.29	2.65	0.01
0.43	0.38	+	_	0.40	+	_	-	_	0.48	0.23	-0.26	_	_	0.75	_	_	-	-	_	-	12	-67.49	165.34	2.00	0.01
0.53	0.07	+	-	0.45	+	-	-	-	0.61	0.52	-0.27	-0.14	-	1.09	-	-	-	-	-	+	14	-64.27	165.47	2.84	0.01
0.53	0.14	+	0.11	0.52	-	-	-	-	0.60	0.56	-0.25	-0.15	-	1.05	-	-	-	-	-	-	13	-65.97	165.52	2.89	0.01
0.29	0.15	+	-	0.44	-	+	-	-	0.52	0.31	-0.25	-	-	0.81	-	-	-	-	-	-	12	-67.60	165.56	2.93	0.01
0.34	0.11	+	-	0.50	+	-	-	-	0.60	0.57	-0.26	-0.14	-	1.05	-	-	-	-	-	-	13	-66.02	165.63	3.00	0.01
0.39	0.47	-	-	0.34	+	+	-	-	0.49	0.29	-0.26	-	-	0.77	+	+	-	-	-	-	15	-62.63	165.69	3.06	0.01
	0.17	+		0.0 1																					
0.27	0.17 0.15	+	-	0.45	-	-	-	-	0.53	0.32	-0.25	-	-0.03	0.81	-	-	-	-	-	-	12	-67.67	165.71	3.07	0.01
					-	- +	-	- -	0.53 0.59 0.50	0.32 0.56 0.24	-0.25 -0.25 -0.13	- -0.14	-0.03 -	0.81 1.04		-		-	-	-	12 13	-67.67 -66.13 -69.29	165.71 165.84 165.86		

Table S15. Model with 90Cl indices (see Table S9 for comparison). Model selection output for the analysis of Delay Index variation excluding Lifespan Effect Index (following on the next page).

	Covariate		Ecolo; facto	-	life	ocial estyle ctors	Life-his facto	-	1	ices refle nge in LR AFR	-			Inte	eracti	ons						Мо	del inform	ation	
(Intercept)	ln(body mass)	sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index 90Cl	Before Variation Index 90Cl	After Variation Index 90Cl	After Variation Index 90Cl : Before Variation Index 90Cl	After Variation Index 90Cl : Choice Index 90Cl	Before Variation Index 90Cl : Choice Index 90Cl	Choice Index 90Cl : helper presence	Choice Index 90Cl : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Likelihood	AICc	Δ AICc	weight
0.41	0.42	+	-	-	-		-	-	0.45	0.27	-0.17	-	-	0.67	-	-	-	-	-	-	10	-70.84	166.00	3.36	0.01
0.32	0.13	+	-	0.49	-	-	-	-	0.59	0.56	-0.25	-0.14	-0.02	1.03	-	-	-	-	-	-	13	-66.22	166.01	3.38	0.01
0.37	0.45	+	-	-	+	+	-	-	0.39	0.19	-	-	-	0.57	+	+	-	-	-	-	13	-66.27	166.13	3.49	0.01
0.43	0.27	+	0.11	0.39	-	-	-	-	0.48	0.23	-	-	-	0.73	-	-	-	-	-	-	11	-69.48	166.24	3.61	0.01
0.43	0.14	+	-	0.38	+	+	-	-	0.56	0.53	-0.26	-0.14	-	0.99	+	+	-	-	-	-	16	-61.11	166.31	3.67	0.01
0.28	0.14	+	-	0.17	+	+	0.12	-	0.38	0.31	-0.28	-	-	0.55	-	-	+	+	-	+	17	-59.22	166.34	3.71	0.01
0.44	0.20	+	-	0.27	+	-	-	-	0.50	0.21	-	-	-	0.77	-	-	-	-	-	+	12	-68.02	166.41	3.78	0.01
0.45	0.24	+	-0.11	-	-	-	-	+	0.49	0.20	-	-	-	0.77	-	-	-	-	-	-	11	-69.59	166.47	3.83	0.01
0.24	0.25	+	-	0.35	-	+	-	-	0.47	0.23	-	-	-	0.71	-	-	-	-	-	-	11	-69.66	166.60	3.97	0.01
0.23	0.24	+	-	0.36	+	-	-	-	0.48	0.23	-	-	-	0.72	-	-	-	-	-	-	11	-69.67	166.62	3.98	0.01
0.66	0.24	+	-	0.55	-	-	-0.29	+	0.62	0.58	-0.20	-0.17	-	1.15	-	-	-	-	-	-	14	-64.87	166.68	4.04	0.01
0.42 0.71	0.04 0.27	+	-	0.47	+	+	-	-	0.56	0.32 0.45	-0.29 -0.12	- -0.12	-	0.87 0.98	-	-	-	-	-	-	13 12	-66.60 -68.25	166.78 166.88	4.15	0.01
	0.27	+ +	-	-	- -	-+	-	+	0.56 0.41	0.45	-0.12 -0.19	-0.12	-	0.98	-	- -	-	-	-	-	12 14	-68.25 -64.97	166.88	4.24 4.25	0.01 0.01
0.45 0.64	0.40	++	-	-	+	++	-	-+	0.41	0.24 0.19	-0.19	-	-	0.62	+ -	+ -	-	-	-	-	14 11	-64.97 -69.80	166.88	4.25 4.25	0.01
0.62	0.29	י +	-	-	-	т -	0.02	+	0.49	0.19	_	_	-	0.76	-	-	_	-	-	-	11	-69.80	166.94	4.23	0.01
0.62	0.28	+	_	_	+	_	-	+	0.49	0.19	_	_	_	0.76	_	_	_	_	_	_	11	-69.83	166.94 166.94	4.30	0.01
0.58							-0.20	+	0.51	0.21	_	-	-	0.81	-	-	_	-	-	-	12	-68.29	166.96	4.32	0.01
	0.28	+	-	0.36	-	-	-0.20	T						- · -											
0.30	0.28 0.29	+ +	-	0.36 0.24	- +	-+	-0.20	-			-	-	-	0.65	+	+	-	-	-	-	14				
0.30 0.46	0.29		-	0.24	- + +	- + +		-	0.43	0.21		- -0.15	-	0.65 1.10	+ -	+ -	-	-	-	-	14 14	-65.01	166.96	4.33	0.01
0.30 0.46 0.58		+	- - -		- + + -		-	-			- -0.29 -0.23	- -0.15 -0.17	- - -	0.65 1.10 1.11	+ - -	+ - -	- - -	- - -	- -	- -	14 14 14				

Table S15 following. Model with 90Cl indices (see Table S9 for comparison). Model selection output for the analysis of Delay Index variation excluding Lifespan Effect Index.

	Covariate		Ecolo fact	-	life	cial style tors	Life-his facto	-		ces refle ge in LR AFR				Inte	ractio	ons					Model information							
(Intercept)	ln(body mass)	Sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index 90CI	Before Variation Index 90CI	After Variation Index 90Cl	After Variation Index 90Cl : Before Variation Index 90Cl	After Variation Index 90Cl : Choice Index 90Cl	Before Variation Index 90Cl : Choice Index 90Cl	Choice Index 90Cl : helper presence	Choice Index 90C1 : family-living	mean lifespan : helper presence	mean lifespan : family-living	nest predation risk : helper presence	nest predation risk : family-living	df	log Lik	AICc	Δ ΑΙCc	weight			
0.55	0.29	+	-0.01	0.62	-	-	-0.34	-	0.61	0.60	-0.23	-0.18	-	1.11	-	-	-	-	-	-	14	-65.15	167.24	4.60	0.00			
0.57	0.29	+	-	0.62	-	-	-0.33	-	0.61	0.60	-0.23	-0.17	-0.01	1.11	-	-	-	-	-	-	14	-65.15	167.24	4.61	0.00			
0.54	0.37	+	-	-	+	+	-	+	0.43	0.17	-	-	-	0.68	+	+	-	-	-	-	14	-65.21	167.35	4.72	0.00			
0.53	0.22	+	-	0.46	-	-	-0.19	+	0.53	0.29	-0.21	-	-	0.86	-	-	-	-	-	-	13	-66.93	167.44	4.81	0.00			
0.26	0.09	+	-	0.16	+	+	0.17	-	0.38	0.31	-0.29	-	-	0.53	-	-	+	+	+	+	18	-57.77	167.45	4.82	0.00			
0.45	0.41	+	-	-	-	-	-	-	0.50	0.45	-0.16	-0.11	-	0.83	-	-	-	-	-	-	11	-70.11	167.49	4.86	0.00			
0.27	0.14	+	-	0.16	+	+	0.14	-	0.39	0.30	-0.26	-	-	0.52	+	-	+	+	-	+	18	-57.81	167.53	4.89	0.00			

Table S15 following. Model with 90Cl indices (see Table S9 for comparison). Model selection output for the analysis of Delay Index variation excluding Lifespan Effect Index.

Model set with Δ AICc \leq 5. N = 36 populations, 34 species.

"+" and "-" indicate the presence or absence of the parameter in the model, respectively. "df" is the degree of freedom. "log Likelihood" is the log likelihood of the model. "AIC_c" represents the Akaike's information criterion corrected for sample size. " Δ AIC_c" is the difference in AIC_c between the focal model and the model with the lowest AIC_c. "weight" represents the relative probability of a model within the full set of models.

	Covariate		1	ogical tors	life	ocial style ctors	Life-his facto	-	Indices reflecting change in LRS or survival with AFR					Inte	eracti	ons			Model information						
(Intercept)	ln(body mass)	sex	latitude	nest predation risk	family-living	helper presence	mean lifespan	chick development mode	Choice Index 90Cl	Before Variation Index 90Cl	After Variation Index 90CI	Lifespan Effect Index	Before Variation Index 90Cl : Choice Index 90Cl	Choice Index 90Cl : helper presence	Choice Index 90Cl : family-living	mean lifespan: helper presence	mean lifespan: family-living	nest predation risk : family-living	df	log Likelihood	AICc	Δ AICc	weight		
0.59	0.22	+	-	0.42	-	-	-	-	0.47	0.32	-0.26	0.44	0.58	-	-	-	-	-	12	-24.05	85.66	0.00	0.06		
0.82	0.53	+	-	-	-	-	-	-	0.42	0.17	-	0.40	0.47	-	-	-	-	-	10	-28.54	85.89	0.23	0.05		
0.88	0.51	+	-	-	-	-	-	-	0.42	0.17	-0.21	0.41	0.48	-	-	-	-	-	11	-26.67	86.34	0.69	0.04		
0.57	0.30	+	-	0.34	-	-	-	-	0.46	0.28	-	0.42	0.54	-	-	-	-	-	11	-27.01	87.02	1.37	0.03		
0.51	0.48	+	-	-	-	-	-	-	0.30	-	-	0.55	-	-	-	-	-	-	8	-32.89	87.10	1.45	0.03		
0.55	0.50	+	-	-	-	-	-	+	0.30	-	-0.31	0.53	-	-	-	-	-	-	10	-29.21	87.22	1.56	0.03		
0.32	0.14	+	-	0.56	-	-	-	+	0.43	0.39	-0.37	0.46	0.47	-	-	-	-	-	13	-22.53	87.60	1.95	0.02		
0.45	0.52	+	-	-	+	+	-	-	0.26	-	-	0.50	-	-	-	-	-	-	10	-29.40	87.61	1.95	0.02		
0.80	0.76	+	0.29	-	+	+	-	-	0.23	-	-	0.51	-	-	-	-	-	-	11	-27.58	88.15	2.50	0.02		
0.57	0.46	+	-	-	-	-	-	-	0.30	-	-0.20	0.56	-	-	-	-	-	-	9	-31.62	88.17	2.52	0.02		
0.49	0.48	+	-	-	+	-	-	-	0.26	-	-	0.56	-	-	-	-	-	-	9	-31.64	88.21	2.55	0.02		
1.00	0.95	+	0.42	-	+	+	-0.18	-	-	-	-	0.49	-	-	-	+	-	-	12	-25.36	88.29	2.63	0.02		
0.50	0.50	+	-	-	+	+	-	-	0.26	-	-0.21	0.51	-	-	-	-	-	-	11	-27.75	88.50	2.84	0.01		
0.39	0.13	+	-	-	-	-	0.36	-	0.28	-	-0.28	0.52	-	-	-	-	-	-	10	-29.97	88.74	3.08	0.01		
0.68	0.99	+	-	-	+	+	-0.49	-	0.24	-	-	0.52	-	-	-	-	-	-	11	-27.95	88.90	3.25	0.01		
0.49	0.51	+	-	-	-	-	-	+	0.30	-	-	0.53	-	-	-	-	-	-	9	-32.08	89.09	3.43	0.01		
0.82	0.55	+	-	-	-	+	-	-	0.44	0.16	-	0.38	0.47	-	-	-	-	-	11	-28.06	89.12	3.46	0.01		
0.55	0.46	+	-	-	+	-	-	-	0.26	-	-0.21	0.57	-	-	-	-	-	-	10	-30.17	89.14	3.49	0.01		
0.40	0.34	+	-	0.24	-	-	-	+	0.32	-	-0.34	0.57	-	-	-	-	-	-	11	-28.28	89.56	3.90	0.01		
0.40	0.30	+	-	-	-	-	0.20	-	0.29	-	-	0.53	-	-	-	-	-	-	9	-32.34	89.60	3.95	0.01		
0.77	0.52	+	-	-	+	-	-	-	0.39	0.18	-	0.41	0.43	-	-	-	-	-	11	-28.33	89.65	4.00	0.01		
0.79	0.52	+	-	-	-	-	-	+	0.40	0.19	-0.27	0.41	0.40	-	-	-	-	-	12	-26.07	89.70	4.04	0.01		
0.59	0.24	+	-	0.41	-	+	-	-	0.48	0.31	-0.25	0.41	0.58	-	-	-	-	-	13	-23.60	89.74	4.08	0.01		
1.01	1.07	+	0.37	-	+	+	-0.29	-	0.14	-	-	0.50	-	+	-	+	-	-	14	-20.90	89.80	4.14	0.01		
0.46	0.46	+	-	-	+	-	-	-	0.20	-	-	0.52	-	-	+	-	-	-	10	-30.51	89.82	4.16	0.01		

Table S16. Model with 90Cl indices (see Table S10 for comparison). Model selection output for the analysis of Delay Index variation including Lifespan Effect Index (following on the next page).

Social Life-history Indices reflecting change in LRS Ecological Model information Covariate lifestyle Interactions factors factors or survival with AFR factors Before Variation Index 90CI : Choice Index 90CI Choice Index 90Cl : helper presence predation risk : family-living 90Cl : family-living lifespan: helper presence mean lifespan: family-living **Before Variation Index 90CI** After Variation Index 90CI chick development mode Lifespan Effect Index nest predation risk 90CI helper presence mean lifespan In(body mass) log Likelihood **Choice Index Choice Index** family-living (Intercept) latitude weight **Δ** AICc mean nest AICc sex đ 0.53 0.01 0.38 0.41 -0.10 0.30 -9 -32.46 89.84 4.19 + _ 0.71 0.43 + -0.10 0.41 0.20 -0.24 0.39 0.46 12 -26.15 89.86 4.20 0.01 _ _ 0.72 0.31 0.20 0.40 0.19 -0.26 0.40 0.43 12 -26.15 89.86 4.20 0.01 + _ 0.74 0.49 -0.05 0.42 0.18 -0.40 0.46 11 -28.4489.87 4.22 0.01 + 0.88 0.53 0.43 0.17 -0.21 0.39 0.48 12 -26.19 89.94 4.28 0.01 + 0.37 0.58 9 -32.52 4.31 0.01 0.40 0.17 0.31 --89.96 + 0.42 11 -28.53 0.01 0.80 0.50 + 0.03 0.17 -0.40 0.46 90.06 4.41 _ 0.82 0.53 0.42 0.17 -0.40 0.46 11 -28.54 90.08 4.43 0.01 + + 0.38 0.36 0.29 -0.24 10 -30.66 0.01 + -0.14 _ 0.54 90.12 4.46 0.51 0.50 0.31 0.53 9 -32.62 90.16 4.51 0.01 _ _ + + 0.71 1.09 + + -0.61 0.20 -0.50 13 -23.81 90.17 4.51 0.01 0.64 0.97 0.48 11 -28.64 90.28 4.62 0.01 + + -0.58 --0.86 0.79 0.35 0.16 _ 0.50 12 -26.36 90.28 4.63 0.01 + _ + 0.83 0.50 + 0.39 0.18 -0.22 0.42 0.44 12 -26.38 90.33 4.67 0.01 0.52 0.19 + -0.05 0.40 0.46 0.32 -0.27 0.43 0.56 13 -23.91 90.37 4.72 0.01 0.57 0.49 0.30 -0.06 -0.56 9 -32.75 90.42 4.76 0.01 + 0.84 0.74 + 0.37 --0.48 10 -30.87 90.53 4.88 0.01 + 0.60 0.23 -0.02 0.47 0.32 -0.25 0.44 13 -24.04 90.62 4.97 0.01 + 0.43 0.58 -0.26 0.44 0.57 13 -24.04 0.59 0.22 + 0.42 + 0.47 0.32 90.64 4.98 0.00 _ 0.52 + + 0.31 -0.31 0.51 11 -28.83 90.65 5.00 0.00 0.54 + -

Table S16 following. Model with 90Cl indices (see Table S10 for comparison). Model selection output for the analysis of Delay Indexvariation including Lifespan Effect Index.

Model set with \triangle AICc \leq 5. N = 22 populations, 21 species.

"+" and "-" indicate the presence or absence of the parameter in the model, respectively. "df" is the degree of freedom. "log Likelihood" is the log likelihood of the model. " AIC_c " represents the Akaike's information criterion corrected for sample size. " ΔAIC_c " is the difference in AIC_c between the focal model and the model with the lowest AIC_c . "weight" represents the relative probability of a model within the full set of models.

Figure S1. Phylogenetic tree for the 34 species studied in this paper (based on the full tree from Jetz et al. 2012; Ericson backbone phylogeny).

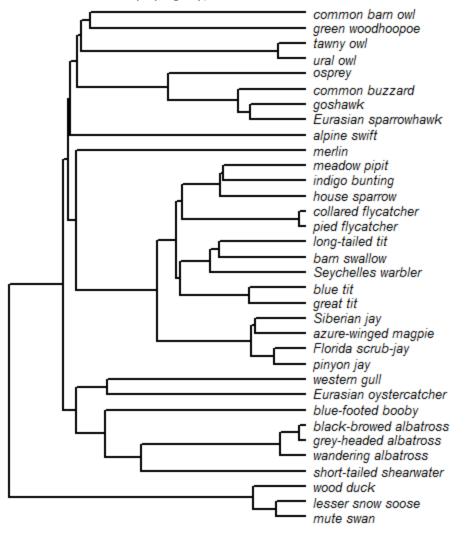


Figure S2. Variation in AFR and consequences for mean reproductive lifespan. Relationship between mean reproductive lifespan (mean lifespan (per AFR classes) minus AFR) and AFR for 22 populations (21 species) used to estimate the Lifespan Effect Index (Table 1); each point represents the mean value for individuals that start to reproduce at a specific AFR. B = both sexes, F = female, M = male.

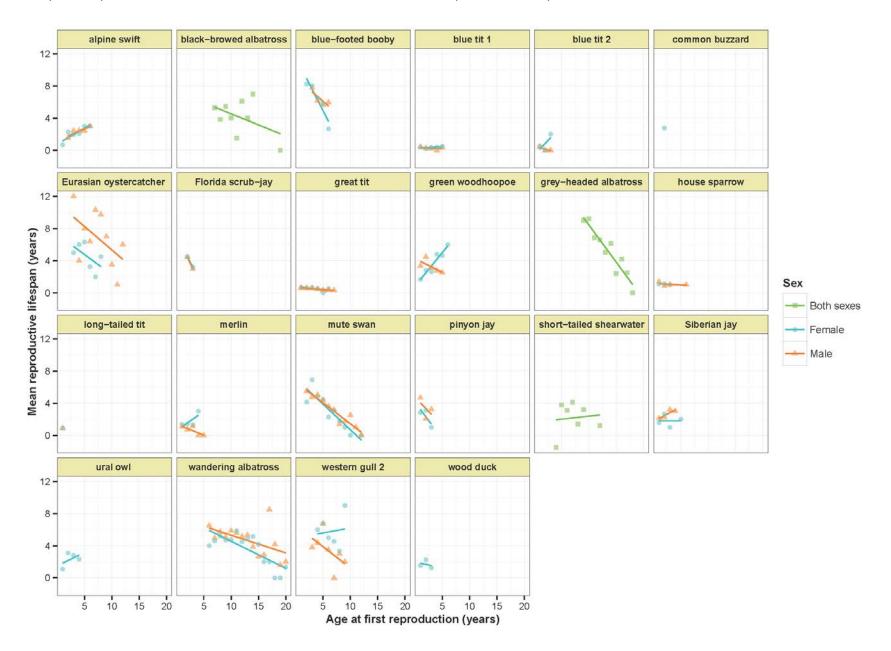
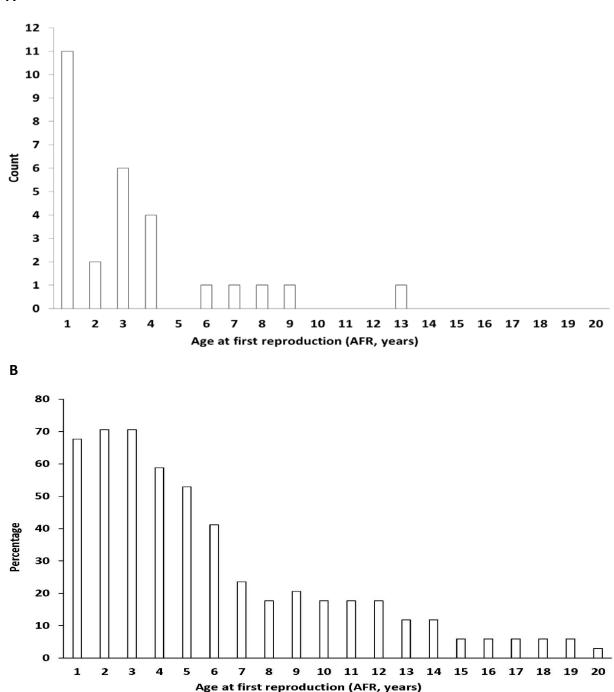
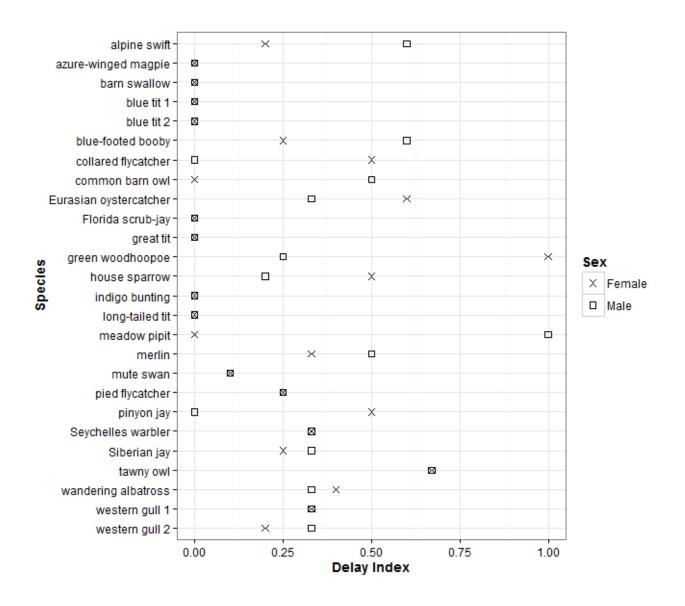


Figure S3. Variation in AFR. (**A**) Number of times the specific AFR corresponded to a species modal AFR (over 28 out of 34 species as we excluded 4 species with only 1 AFR age class and 2 species for which the sample size per AFR age class was missing). (**B**) Frequency of observation of a specific AFR age class across all 34 species (an AFR age-class was counted as being observed within a population when at least one individual initiated reproduction at the focal AFR – e.g. a values of about 20% for an AFR of 9 means that about 7 species (20% of 34) had individuals that initiated their reproduction at age 9).



Α

Figure S4. Sex differences in the Delay Index for the 26 populations (24 species) for which we had separate data for males (M, square symbols) and females (F, cross symbols). A number after the species indicates separate studies.



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