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Climate warming and temporal variation in

2 reproductive strategies in the endangered meadow

3 viper

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- 20 editorial advice.
- 21 **Running headline:** Climate warming and reproductive strategies in a viper

22 Summary

23 Anthropogenic climate change poses a significant threat to species on the brink of extinction. 24 Many non-avian reptiles are endangered, but uncovering their vulnerability to climate 25 warming is challenging because this requires analysing the climate sensitivity of different life 26 stages and modelling population growth rates. Such efforts are currently hampered by a lack 27 of long-term life history data. In this study, we used over three decades of mark-recapture data 28 from a natural population of the endangered meadow viper (Vipera ursinii ursinii) to unravel 29 the patterns of temporal variation in reproductive traits, the local climatic determinants of 30 inter-annual variation in reproduction and the potential buffering effects of life cycle on 31 population growth rate. We found significant inter-annual variation in body growth, gestation 32 length, post-parturition body condition, clutch success and offspring traits at birth, while 33 reproductive effort showed little temporal variation. Temperature during gestation was the 34 most critical factor, reducing gestation length and increasing both clutch success and post-35 parturition body condition. In contrast, neither air humidity nor global radiation affected 36 reproductive outcomes. This population had a negative growth rate with minimal temporal 37 variation, indicating a rapid decline largely independent of climatic conditions. Overall, the 38 viper's life history traits appeared to be buffered against temporal variation in climatic 39 conditions, with this declining population potentially benefiting on the short-term from rising 40 local temperatures.

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42 Keywords: reproduction, body growth, temperature, rainfall, gestation.

44 Introduction

45 Habitat loss, overexploitation of wild populations for food or resources, and introduced alien 46 species have pushed many species to the brink of extinction (Ceballos et al. 2020). 47 Anthropogenic climate change is an aggravating extinction threat and thus plays a major role 48 in the current biodiversity crisis (McLaughlin et al. 2002; Urban 2015). To improve our understanding of climate-driven extinction processes, a consensus has emerged on the need 49 50 for integrated assessments of climate vulnerability (Williams et al. 2008; Huey et al. 2012). In 51 this conceptual framework, extinction risk depends not only on exposure to climate change, 52 but also on the sensitivity and resilience of the organism. Studying the extinction risk of 53 species with complex life cycles, such as many iteroparous organisms or those living in highly 54 seasonal environments, requires an understanding of the effects of climate change at different 55 life stages, and the modelling of population growth rates (reviewed in Brodie et al. 2013). 56 In this context, long time series of demographic data from individually marked 57 individuals are needed to decipher direct and delayed effects of climatic conditions, to account 58 for potential compensatory responses across the life history, and to quantify specific effects of 59 extreme weather events that rarely occur in nature (Williams et al. 2008; Brodie et al. 2013). 60 Integrated climate vulnerability assessments exist for a few plants and animals, but are scarce 61 for non-avian reptiles (turtles, lizards, snakes, and crocodiles) due to the paucity of long-term 62 data sets for these organisms (Le Galliard et al. 2012; Böhm et al. 2013; Winter et al. 2016). 63 Furthermore, non-avian reptiles are at high risk of extinction compared to other tetrapods, 64 with more than 20% of existing species being currently threatened, but the specific threats 65 posed by climate change remain highly uncertain for most of these organisms (Gibbons et al. 66 2000; Falaschi et al. 2019; Cox et al. 2022). Thus, long-term studies of reptile population 67 dynamics are needed to improve our understanding of the patterns and processes of their 68 declines (Diele-Viegas and Rocha 2018; Diele-Viegas et al. 2020).

69 Non-avian reptiles are highly sensitive to the direct effects of global warming due to 70 their behavioural thermoregulatory strategies (i.e., ectothermy). Direct, negative effects of 71 climate change may include increased mortality risks due to overheating (e.g., Kingsolver et 72 al. 2013), sublethal effects due to reduced basking and foraging activities (e.g., Sinervo et al. 73 2010), disruption of energy balance (reviewed in Rutschmann et al. 2024), or increased 74 likelihood of physiological dehydration (e.g., Kearney and Porter 2004). Such deleterious 75 effects can alter early life stages, whose survival depends strongly on abiotic conditions, but 76 also adult life stages, especially in viviparous species where females must optimally regulate 77 their body temperature during a long gestation period (Shine 2003; Le Galliard et al. 2012). 78 However, the cumulative effects of these stresses on population growth and viability are 79 uncertain, as non-avian reptiles have also developed strategies to buffer the negative effects 80 of climate variability (Le Galliard et al. 2012; Cox et al. 2022). For example, some are 81 efficient thermoregulators and exploit habitat heterogeneity to maintain homeostasis, even 82 under changing environmental conditions (Huey et al. 2012; Sunday et al. 2014). On the other 83 hand, many iteroparous reptiles have low and flexible metabolic requirements and can cope 84 with prolonged periods of starvation or chronic water deprivation (Shine 2005). Some are 85 capital breeders, accumulating energy for several months or even years before breeding (Bonnet et al. 1998; Shine 2003). For these species, we expect small environmental 86 87 fluctuations in cumulative lifetime reproductive success, even under changing climatic 88 conditions. Unfortunately, studies quantifying the temporal variability of life history traits and 89 population growth rates are rare in reptiles (but see Miller et al. 2011; Rodríguez-Caro et al. 90 2021; Warret Rodrigues et al. 2021).

Climate change may also have indirect effects on non-avian reptile populations via its
impact on the quantity and distribution of food resources, or the abundance of predators (see
Diele-Viegas and Rocha 2018 for a review). The trophic effects of climatic conditions have

94 been well studied in ecosystems characterized by resource pulses; for example, high rainfall 95 regimes in certain arid or tropical environments favour greater prey abundance and thus have 96 positive effects on the growth, reproduction, or survival of certain snake species (e.g., Ujvari 97 et al. 2010; Meik et al. 2024). In temperate regions, climate change may also alter the prey 98 community, with potentially variable effects on reptile demography between studies (Diele-99 Viegas and Rocha 2018). For example, in a viper population in Italy, whose phenology is 100 highly dependent on ambient temperature, prey availability reflects local habitat variation 101 rather than climatic conditions (Rugiero et al. 2012, 2013). In contrast, in another viper 102 population in Sweden, an extreme summer climatic event caused a potential decline in 103 resources that was associated with a loss of body condition and a reduction in future adult 104 survival (Madsen et al. 2023). These trophic effects of climate change are likely to be more 105 pronounced in reptiles with specialized diets and inflexible foraging behaviour (Ujvari et al. 106 2011; Diele-Viegas and Rocha 2018).

107 Among non-avian reptiles, vipers rank among the most endangered groups, with 16% of 108 species threatened compared to 8% of all reptiles' species (Böhm et al. 2013; Maritz et al. 109 2016; Cox et al. 2022). Many snake populations are declining globally, primarily due to 110 habitat loss or reduced prey availability (Reading et al. 2010; Luiselli et al. 2011; Böhm et al. 111 2013). However, changes in climatic conditions can also drive long-term population declines 112 (e.g., Shine and Mason 2004; Pomara et al. 2014; Madsen et al. 2023). Snakes exhibit 113 ecological traits that are sensitive to climate change, with higher temperatures generally 114 enhancing activity levels, accelerating gestation, and improving the quality of reproduction in 115 temperate species (Lourdais et al. 2004; Moreno-Rueda et al. 2009; Le Galliard et al. 2012; 116 Rugiero et al. 2013). On the other hand, winter and summer climatic conditions can have 117 variable effects on annual survival rates (Altwegg et al. 2005; Pomara et al. 2014; Elmberg et 118 al. 2024). Extreme climate events, especially those leading to water shortage, are also likely to

disrupt resource availability and reproductive success (Dezetter et al. 2021; Holden et al.

120 2022; Madsen et al. 2023).

121 Here, we analysed over three decades of mark-recapture data from a population of the 122 meadow viper (Vipera ursinii ursinii, Bonaparte 1835) to uncover patterns of temporal 123 variation in reproductive traits, to identify the local climatic factors influencing these 124 variations, and assess the potential buffering effects of life cycle on population growth rates. 125 Vipera u. ursinii is a small viper with a fragmented distribution across medium-altitude 126 mountain ranges in the northern Mediterranean basin. This species is listed on the IUCN Red 127 List of threatened European species, although climate change is not currently identified as a 128 primary threat (Ferchaud et al. 2011). Previous research on our study population has revealed 129 that the meadow viper has a slow life history with a continuous growth throughout life, a 130 sexual maturation at 3-5 years of age, a biennial breeding, and a high and stable annual 131 subadult and adult survival (Baron et al., 1996, 2013; Tully et al., 2020). In our study 132 population, the vipers primarily feed on large grasshoppers, which represents up to 99% of the 133 diet during summer (Baron 1992). Although growth rates and some reproductive traits 134 fluctuate from year to year, it is unclear whether this variability is linked to climatic 135 conditions, or to what extent life history traits variability contribute to temporal variation in 136 the population growth rate.

In this study, we quantified the effects of climatic variables related to the most relevant thermal and water constraints. Since the meadow viper is a capital breeder, alternating between breeding and non-breeding years (Baron et al. 2013; Tully et al. 2020), we tested for the effects of climatic condition during gestation, during the mating period prior to gestation, and during the preceding non-reproductive year. We next parametrized a finite stochastic integral projection model (Easterling et al. 2000; Vindenes et al. 2011; Jaffré and Le Galliard 2016) to (1) quantify the demographic and environmental variances in population growth rate,

144 and (2) identify the vital rates most critical for population growth. Our analysis revealed

145 distinct patterns of climate-reproduction relationships, but minimal environmental variation in

- 146 population growth rate. Our findings suggest limited potential for direct effects of climate
- 147 warming on the vulnerability of this population.

148 Materials and methods

149 Study species and site

150 The meadow viper *V. u. ursinii* is a small insectivorous viper inhabiting open and dry

151 calcareous grasslands between 900 and 2 200 m a.s.l. in the northern Mediterranean basin.

152 This study was carried out in Mont Ventoux population (1430 m altitude, 44°18'N, 5°26'E), a

153 population occupying an area of 100 ha that is isolated from the nearest populations 45 km

away. A four hectares study area has been monitored continuously from 1979 until 2014,

155 when monitoring was halted due to lack of support and conflicts with local landowners. The

area is divided into two contiguous study sites of approximately two hectares each, including

157 a north-facing and mesophilic hillside (habitat A), and a south-facing and xerophilic hillside

158 (habitat B). Small home ranges averaging 0.09 ha (Baron et al. 1996), together with the

159 presence of a 10 m wide asphalt road separating the two sites, make movements of snakes

160 between these two sites extremely rare (only two such movements were reported during this

161 study). In this population, females become sexually mature at approximately 4-6 years of age.

162 Adult males and adult females emerge around mid-April and early May respectively.

163 Vitellogenesis begins after hibernation and lasts 4 to 8 weeks. Mating occurs during the last

164 two weeks of May, soon followed by ovulation and fertilization, which occurs during the first

165 week of June. Parturition takes place between mid-August and mid-September, depending on

166 weather conditions. Hibernation begins in October.

167 **Demographic data collection**

168 Snakes were individually marked by scale clipping upon their first capture during 56 primary 169 capture sessions between 1979 and 2014 (n=1346 captures, mean duration per session of 10 170 days). Although scale clipping is highly reliable, we also recorded ventral and head scales to 171 confirm the ventral scale clipping pattern. Primary capture sessions were carried out by 1-3 172 people for 1-2 weeks twice a year, including one session during the mating season and before 173 ovulation, and one session before the end of gestation. All spotted animals were captured, 174 identified and measured (snout-vent length, SVL to the nearest mm and body mass to the 175 nearest dg). Sexual maturity and reproductive status (non-reproductive, reproductive: post-176 ovulation, gravid or post-partum) of each female was assessed. During the second capture 177 session, ova were counted by palpation of the abdominal cavity to provide an accurate estimate of litter size (Baron et al. 2013). Snakes were released at their capture site in the 178 179 evening of the day of capture, or after parturition for gravid females (see below). All 180 applicable institutional and/or national guidelines for animal care and use were followed. 181 In 1983-1988 and from 1994 onwards, we obtained data on reproductive output by 182 keeping gravid females in the laboratory until they gave birth (498 offspring, 162 litters, 109 183 mothers, 28 years). Gravid females were kept in individual rearing boxes ($350 \times 180 \times 210$ 184 mm) with a shelter, damp soil, free access to water and a heat source to allow the snakes to 185 thermoregulate. Body mass was measured daily. Immediately after parturition, the number of 186 undeveloped ova, dead embryos, stillborns and healthy offspring were counted to calculate 187 litter size and litter success (all variables are precisely defined in Baron et al. 2013). On 188 average, females were kept in the laboratory for 15 days (\pm 10.7 SD, range = 0-46) prior to 189 parturition, a short period of captivity that should not influence reproductive parameters. The 190 gestation period was defined as the time between the date of ovulation (assumed to start on 10 191 June based on pers. obs.) and the date of parturition. Newborns were measured for SVL and

body mass, and sexed by counting sub-caudal scales. Mothers were released with their
offspring at the maternal capture site on average 11 days (± 7 SD) after parturition. Data on

194 survival and growth during the juvenile stage and data on intermittent reproduction and

195 longevity have been reported elsewhere (Baron et al. 2010a, 2013; Tully et al. 2020).

196 Climate data collection

197 Temperature and rainfall data were recorded since June 1992 by Météo France at a non-198 automated weather station at the study site (Beaumont-Mont Serein, 1445 m a.s.l., 199 44°10'48"N, 5°15'18"E). To reconstruct time series of weather data during the entire time 200 period of the census (1980-2014), we gathered temperature and rainfall data from an 201 automated meteorological station located 30 km south-west of the study site (Carpentras, 99 202 m a.s.l., 44°05'00"N, 5°03'30"E), and temperature, rainfall, global radiation and specific 203 humidity data calculated for the study site by the Safran model of Météo France (Quintana-204 Seguí et al. 2008). The Carpentras station provides high quality and long term meteorological 205 data on temperature and precipitation, but is located at a lower altitude than the study site. The 206 Safran model is a gauge-based analytical meteorological system that provides reconstructed 207 and robust weather data through interpolation methods at 8 km resolution in France for the period 1958-2014 (Vidal et al. 2010). We obtained daily data on precipitation (including 208 209 snowfall), minimum and maximum air temperature, specific humidity, wind speed and global 210 radiation for a grid cell centred on the study site. For this study, we calculated daily maximum 211 temperature (°C), daily cumulative rainfall (mm), daily cumulative global radiation (J/cm²) 212 and daily mean of atmospheric specific humidity (g/kg) as climate variables (Appendix A). 213 We chose these variables because they are relevant to the thermal and water biology of 214 reptiles such as the meadow viper (Lourdais et al. 2004; Guillon et al. 2009; Le Galliard et al. 215 2012), and may correlate with environmental constraints on basking time (low daily 216 maximum temperatures and low global radiation) and environmental constraints on

217 evaporative water loss and water uptake (limited rainfall and low air specific humidity). For 218 each climate variable and each study year, we calculated stage-specific weather data during 219 the mating period (the three first weeks of May), the average gestation period (from the first 220 week of June to the first week of September), and the non-reproductive period of the previous 221 year (from the second week of June to the last week of September of the previous year). The 222 non-reproductive period of the previous year was included to account for intermittent biennial 223 reproduction and the capital breeding strategy of the species (Baron et al. 2013). In addition, 224 we calculated the temperature conditions experienced during three consecutive gestation 225 stages, namely early gestation (first week of June to first week of July), mid-gestation (first 226 week of July to first week of August) and late gestation (first week of August to first week of 227 September).

228 Statistical analyses

229 We first used mixed-effects models to investigate temporal variation (between years) and 230 spatial variation (between sites A and B) in gestation length, total clutch size and clutch 231 success (percentage of viable offspring in the clutch) controlling for body size, variation in 232 clutch mass (including water and residual tissue from amnion and allantois) and variation in 233 post-partum body condition (log-transformed body mass after parturition, controlling for log-234 transformed body size). We analysed clutch mass including post-partum body mass as a 235 covariate, which amounts to examining variation in "relative clutch mas" (Van Dyke and 236 Beaupre 2011). We also calculated mass change and body size growth of females during their 237 non-reproductive years (57 observations, 35 females), as we have previously shown that 238 females stop growing and lose significant body mass during the reproductive years (Baron et 239 al. 2013). Models for for body mass change and body size growth included initial body mass 240 and initial body size as covariates in the models, respectively, to account for the slowing of 241 growth with age (Baron et al. 2010b). Inter-individual differences were also tested prior to full

242 model implementation, but we found that these were negligible compared to inter-annual 243 differences so we did not include them in our final models. We also analysed spatial and 244 temporal variation in offspring characteristics, including mass and body condition at birth. 245 Body condition at birth was defined as the residual of a linear regression of body mass against 246 SVL. Body mass was preferred to SVL for analyses of variation in offspring size because (1) 247 offspring mass can be measured more accurately than SVL, (2) mass better reflects maternal 248 energy allocation to the offspring than SVL, and (3) mass at birth is often used in studies of 249 temporal variation in offspring fitness traits in reptiles (Baron et al. 2010a). 250 In all cases, we followed the methodology of Pinheiro and Bates (2000) for general 251 linear models (normally distributed data) and of Bolker et al. (2009) for generalized linear 252 models (binomially distributed data). In all models, habitat (site A vs. B) was included as a 253 fixed effect and year was included as a random effect. We used a random intercept model, 254 which assumes that years vary in the mean values of each response trait. We also checked for 255 temporal autocorrelation by calculating the empirical autocorrelation function for the within-256 group residuals of the fitted models and found that temporal autocorrelation was negligible. 257 For offspring traits, we included the random effect of litter identity, nested within birth year 258 and habitat effects. Mixed models were fitted using the *lme* package (Pinheiro et al. 2022) in 259 R 3.2.4 software (R Core Team 2020). Model assumptions (normality, homoscedasticity) 260 and/or overdispersion) were checked and met in all cases.

We first used these models to quantify variation between years and maternal identity, controlling for fixed effects of critical covariates (body size and habitat). Fixed effects were tested with conditional F-tests based on REML and random effects were tested with likelihood ratio tests. Initial models included all factors and model selection was performed backwards by removing non-significant terms until a minimum adequate model was obtained (all factors p > 0.05 removed). From the minimum adequate model, we calculated the ratio of annual

variation to the sum of annual and residual variation, hereafter referred to as relative temporalvariation (RTV).

269 In a second step, we used the minimum adequate model as a starting point to search for 270 the most influential weather covariates at each life stage. Following our investigation of all 271 climate variables (see below), we tested for effects of temperature, global radiation and 272 specific humidity calculated during the mating, gestation and non-reproductive periods on 273 reproductive traits with a significant inter-annual variation (see below). We compared all 274 models with additive effects of each covariate and therefore did not test for interactions or 275 quadratic effects. We however checked graphically for the absence of quadratic effects of 276 climatic conditions (results not shown). We compared the independent effects of temperature, 277 global radiation and specific humidity to determine which weather variable was the best 278 predictor of inter-annual variation. Where climatic conditions during gestation were retained 279 in the best model, we next tested more specifically for the effects climatic conditions 280 calculated during early (first month), mid (second month) or late (third month) gestation 281 period. Models were compared by Akaike Information Criterion corrected for small sample 282 size (AICc), Akaike's weight (wi) and evidence ratio (ER) using the MuMIn (Multiple-Model 283 Inference) package (MuMIn, Bartoń 2022). In addition, we calculated the marginal pseudo-R2 284 (R2m) and the conditional pseudo-R2 (R2c) for each mixed-effects model using the 285 *r.squaredGLMM* function of the *MuMin* package (Nakagawa and Schielzeth 2013). The best 286 set of models included all models within a cumulative Akaike's weight < 0.95, and the best 287 models were considered to be those with an AICc difference with the best model < 2288 (Burnham and Anderson 1998, see Appendix B for a list of model selection tables). Relative 289 variable importance was calculated as the sum of wi of all models including the variable, and 290 parameter estimates were estimated using a model averaging procedure (conditional average). 291 Results are presented as mean \pm SE unless otherwise stated.

292 **Results**

293 Climate variation

294 Supplementary information on temporal variation and correlation patterns between climate 295 variables are provided in Appendix A. The weather at the study site showed marked seasonal 296 variation, with harsh thermal conditions for reptile activity from November to April (daily 297 maximum temperatures around and below 5°C and daily minimum temperatures below 0°C), 298 a short summer season from June to August (daily maximum temperatures above 15°C and 299 daily minimum temperatures above 10°C), and heavy rainfall or snowfall from September to 300 May (cumulative monthly rainfall or snowfall greater than 100 mm). The temperature and 301 rainfall data collected at the study site since 1992 were highly correlated with those of the 302 Safran model available on a long-term basis, allowing time series of temperature and rainfall 303 conditions to be reconstructed for the entire duration of the population survey. There was a 304 weak pattern of increasing daily maximum temperatures in late spring and early summer over 305 time during the study, but not for average summer temperatures (see Appendix A and Figure 306 A1). Global radiation increased significantly over time during the survey (Figure A1). 307 Cumulative rainfall and air specific humidity showed no temporal trend. Independent of these 308 trends, weather data showed strong inter-annual variation at all reproductive stages. As 309 rainfall at the study site was less well predicted by the Safran data, and was also negatively 310 correlated with global radiation and positively correlated with air specific humidity, we did 311 not use rainfall data in our statistical models of life history variation in vipers.

312 Life history variation

Similar to previous findings (Baron et al. 2013; Tully et al. 2020), certain reproductive traits
and body growth were significantly influenced by maternal body size and habitat (Table 1).
Larger females produced larger litters, exhibited higher clutch success, and gave birth to

316 larger offspring. Additionally, females from the xerophilic habitat (Habitat B) displayed 317 similar size-independent fecundity and reproductive effort to those of females from the 318 mesophilic habitat (Habitat A). However, they exhibited lower post-parturition body 319 condition, reduced offspring condition, and smaller body size growth and annual change than 320 females from Habitat A. Moreover, we detected significant inter-annual variation in all 321 demographic traits, except for total clutch size (Table 1). RTV was generally low for 322 reproductive effort, where it was largely driven by a few years with unusually low 323 reproductive effort and also for offspring mass. Intermediate levels were observed for clutch 324 success, post-parturition body condition, offspring condition and body size growth. The 325 highest RTV was recorded for gestation length and annual changes in body mass during non-326 reproductive years (detailed results provided in Table 1).

327 Climate effects on reproduction and growth

328 Our model selection procedure included 17 candidate models and one null model (the 329 minimum adequate models shown in Table 1, see Appendix B for the full list of all models). 330 We excluded from the set of candidate models some models with additive effects of global 331 radiation variables due to strong collinearity between the variables (see Appendix A). The 332 combined outcomes from our model selection procedures for gestation length (Table B1), 333 clutch success (Table B2), relative clutch mass (Table B3), post-parturition body condition 334 (Table B4) and offspring mass and condition (Tables B5 and B6) showed very small effects of 335 specific humidity, moderate effects of global radiation and significant positive effects of air 336 temperature.

337 The gestation length shortened significantly with increasing average temperatures

during both the mating period (-1.46 days/ $^{\circ}$ C ± 0.48, z = 2.92, P = 0.003) and gestation (-2.76

days/°C \pm 0.96, z = 2.73, P = 0.006; Figure 1, Table B1), with relative importance (*w*) of 0.94

and 0.93 respectively. It also tended to shorten with hotter thermal conditions during the

previous non-reproductive season (w = 0.46). The best model, which included average temperature during gestation (Table B1, wi = 0.49), outperformed the three other gestationstage dependent models. The second-best model included effects of temperature during early gestation (wi = 0.45). Upon re-analysing the data excluding the coldest summer season, which saw a high number of clutch failures (Figure 1A), we obtained similar results, with even stronger effects of temperatures (gestation: -3.75 days/°C ± 0.79, z = 4.5, P < 0.0001; mating: -1.57 days/°C ± 0.38, z = 3.92, P < 0.0001).

Climatic conditions explained little variation in relative clutch mass, with all covariates 348 349 showing relative importance below 0.25 (Table B2). The most influential covariate was global 350 radiation during the previous summer season, and it negatively affected relative clutch mass (-351 0.009 ± 0.004 , z = 1.97, P = 0.05, result not shown). In sharp contrast, thermal conditions 352 were major drivers of inter-annual variation in post-parturition body condition (Table B3). 353 Average temperature during gestation was the most influential variable (w = 0.66), followed 354 by average temperature during the previous year (positive effect, w = 0.48) and during the 355 mating period (negative effect, w = 0.45). Post-parturition body condition increased with 356 rising average temperatures during gestation (0.031 \pm 0.014 g/°C, z = 2.1, P = 0.04; Figure 357 2A). Based on additional analyses, we found that late gestation thermal conditions were the 358 primary driver of this pattern (wi = 0.65; 0.024 ± 0.009 g/°C, z = 2.4, P = 0.01). 359 Clutch success increased on average with higher mean temperature during gestation 360 (logit regression: 0.45 ± 0.21 , z = 2.1, P = 0.04; Figure 2B, Table B4, w = 0.51), and was 361 moderately influenced by temperature and humidity during the previous non-reproductive 362 season (w = 0.34 and w = 0.21, respectively). Additional analyses indicated that the model 363 with average temperature during gestation fitted equally well the data than a model with

average temperature during mid-gestation (wi = 0.49 and wi = 0.45, respectively). Climatic

365 conditions did not significantly influence offspring mass at birth (Table B5, all w < 0.22).

However, both mean global radiation (w = 0.54, 0.0006 ± 0.0002 , z = 2.9, P = 0.004) and temperature during gestation (w = 0.24, 0.05 ± 0.02 , z = 2.1, P = 0.04) were positively correlated with offspring condition at birth. Lastly, annual changes in body mass and body growth during the non-reproductive years were unaffected by climatic conditions during the active summer season (all w < 0.23, not shown).

371 Population growth analysis

372 We fitted a finite stochastic integral projection model (IPM) to the meadow viper life cycle 373 and analysed the stochastic IPM using the methodology described in Jaffré and Le Galliard 374 (2016, see Appendix C). The deterministic IPM predicted a substantial annual population decline of approximately 12% (*Section 2019*), consistent with previous findings using a matrix 375 376 population model (Baron et al. 1996). Components of stochastic growth included a significant demographic variance () and a very weak environmental variance (377 378), indicating that the iteroparous life cycle with a constant adult survival and breeding 379 frequency buffered the population against temporal variation in growth, juvenile survival and 380 reproduction. In addition, an analysis of the elasticity of the deterministic growth rate to 381 changes in the kernel function of the IPM revealed (1) very low sensitivity to changes in adult 382 reproduction, (2) strong sensitivity to the survival and growth of juveniles and immature 383 females, and (3) moderate sensitivity to the survival and growth of older sexually mature 384 individuals (Figure 3A). Using a realistic starting population size of 100 vipers, the stochastic 385 model predicted a median extinction time of 25–30 years for this population, with complete 386 extirpation of the population likely within less than a century (Figure 3B). Given the weak 387 contribution of reproduction to growth rate, climate warming effects on reproductive traits is 388 unlikely to significantly alter this extinction trajectory.

389 **Discussion**

390 Our analysis complements previous research on body growth rate strategies, survival, 391 intermittent reproduction, and ageing in the same meadow viper population (Baron et al. 392 2010a, b, 2013; Tully et al. 2020). In particular, our findings reveal that fecundity and 393 reproductive effort are largely buffered against environmental variation and are primarily 394 dictated by female body size. Female fecundity is strongly constrained by body size and 395 exhibit little to no variability among years. Similarly, reproductive effort, defined as the 396 relative mass lost during gestation (Bonnet et al. 2002; Van Dyke and Beaupre 2011), shows 397 minimal inter-annual variation, except for one year marked by a notably poor reproduction 398 without any clear climatic explanation (Table 1). These results contrast with the pronounced 399 vear-to-year variation in reproductive effort and frequency seen in other snake species, which 400 often occurs in response to changes in food availability. For example, some snakes feed on 401 prey whose populations fluctuate dramatically due to changes in rainfall patterns, leading to 402 pronounced reproductive variability in those species (Madsen and Shine 1999; Lourdais et al. 403 2002; Ujvari et al. 2010). This discrepancy likely stems from the meadow viper's unique 404 semi-insectivorous diet, which consists almost exclusively of crickets and grasshoppers. At 405 our study site, these insects are both diversified and abundant during the summer activity 406 season (Baron 1992), indicating that food resources are unlikely to be limiting factors and this 407 stable supply of food may contribute to the low variability in reproductive traits observed in 408 this population.

In contrast to fecundity and reproductive effort, we observed a strong temporal variability in annual growth rates during non-reproductive years, as well as in traits related to gestation speed, body condition, and reproduction quality. These variations were either unrelated to climatic conditions even during extreme climate years, such as body growth of adult females and body mass of young at birth, or were linearly correlated with ambient

414 temperature rather than air humidity (which indicates desiccation risk due to evaporative 415 water loss) or global radiation (which indicates both thermoregulatory conditions and 416 rainfall). In general, warmer ambient conditions are more favourable for reproduction by 417 accelerating gestation, improving post-parturition condition, and ensuring better offspring 418 condition at birth.

419 The flexibility of gestation length and post-partum body condition, which is driven by 420 warmer temperatures, is not surprising but demonstrates that reproductive traits can be 421 flexible without significantly affecting reproductive effort or frequency. This should be 422 understood in the light of the meadow viper's bioenergetic strategy of biennial reproduction 423 and resource capitalization (Bonnet et al. 1998; Baron et al. 2013; Huang et al. 2013). Females 424 alternate between breeding years - marked by a halt in structural growth and depletion of 425 body reserves - and non-breeding years, during which growth resumes and body condition is 426 restored to a threshold necessary for future reproduction (Baron et al. 2013). Our findings 427 suggest that warmer ambient temperatures during gestation accelerate reserve capitalization dynamics but do not alter the biennial threshold for reproduction (Madsen and Shine 1999; 428 429 Baron et al. 2013; Huang et al. 2013; Capula et al. 2016). These effects on reserve dynamics 430 are unlikely to influence future survival, as survival is not linked to post-partum body 431 condition (Baron et al. 2013).

In contrast, lowland populations of the asp viper (*Vipera aspis*) exhibit a more semelparous reproductive strategy, where higher gestation temperatures result in poorer body condition by the end of gestation (Lourdais et al. 2002, 2004). One explanation for this difference is that meadow vipers are able to feed during gestation, unlike carnivorous vipers that are anorexic due to the challenge of consuming large preys while gravid (Bonnet et al. 2002; Baron et al. 2013). Thus, we hypothesize that warmer thermal conditions during gestation may favour maternal feeding in meadow vipers, enabling them to recoup the energy

expenditure of viviparity more efficiently. Our analysis further suggests that thermal
conditions toward the end of gestation, during late summer, are particularly important for
maintaining female energy balance and reserve dynamics.

442 The impact of ambient temperatures on the speed of gestation reflects a general effect of 443 thermal conditions on the rate of biological processes in ectotherms, even including years of 444 heat waves that might have slowed gestation by altering daily activities during the hottest 445 times of the day. Such phenological changes are common in many temperate non-avian 446 reptiles (Lourdais et al. 2004; Moreno-Rueda et al. 2009; Rugiero et al. 2013; Diele-Viegas 447 and Rocha 2018). Here, it corresponded to a shift of about three weeks between the coldest 448 and warmest years, a pattern very similar to that observed in the asp viper (Lourdais et al. 449 2004; Lorioux et al. 2013). In this congeneric species, thermal conditions during mid-450 gestation are more strongly correlated with gestation length, whereas our study suggests that it 451 is the average temperature over the entire gestation that is most important. Thus, generally 452 warmer summer conditions, in early, mid or late gestation, will accelerate gestation similarly. 453 The demographic consequences of shifts in reproductive phenology are likely to be small in 454 meadow vipers, as the date of parturition does not affect the future maternal or neonate 455 survival, growth and sexual maturation (Baron et al. 2010a, 2013).

456 A notable demographic effect of climatic conditions is a positive correlation between 457 average temperatures during gestation and reproductive success, measured by the proportion 458 of viable neonates in the litter. This trend holds even when accounting for data from the 459 extreme heat wave of 2003, when summer temperatures were unusually high and could have 460 imposed sublethal stress on reproduction. Overall, reproductive success is typically high in 461 this species, but cooler years are associated with an increased frequency of developmental 462 failures and prolonged, incomplete gestations. This reflects the importance of optimal 463 thermoregulation during gestation for embryo development and viability in vipers (Lorioux et

464 al. 2013). Altogether, we found that some reproductive traits in this population are currently 465 constrained by cold thermal conditions. Thus, assuming other factors remain constant, climate 466 warming is likely to provide short-term benefits to this population, a pattern that may apply 467 more generally to high-altitude reptiles, whose energy budgets are constrained by colder and 468 shorter activity seasons (Huang et al. 2013). Comparative data and mechanistic models could 469 be used to determine whether this trend is consistent across the species' range. For example, 470 similar studies could examine populations in southern regions or at lower elevations, which 471 are closer to the species' upper thermal limits, to see if they respond differently to climate 472 warming.

473 We integrated the effects of the different vital rates and their temporal variability on 474 long-run growth rate and population dynamics using the Integral Projection Model framework 475 (Easterling et al. 2000; Ellner and Rees 2006; Vindenes et al. 2011). The motivation for 476 building this model is the existence of a high variability in body size, with pervasive effects of 477 individual size on growth, sexual maturation, fecundity, or the size of newborns at birth. Our 478 analysis of an individual-based version of the model allows for both demographic fluctuations 479 generated by random, identically and independently distributed temporal variability in each 480 vital rate (i.e., environmental stochasticity) and random inter-individual variability in the 481 demographic processes of growth, survival, and reproduction (i.e., demographic stochasticity). 482 This shows that temporal fluctuations in the stochastic growth rate are negligible compared to 483 random inter-individual demographic fluctuations in the meadow viper. Assuming that the 484 average stochastic growth rate scales like (Vindenes et al. 2011; 485 Jaffré and Le Galliard 2016), the population size at which the contribution of demographic 486 variance equals contribution of environmental variance would be approximately 1,000 487 females in this population, well above the estimated population size of a few hundred vipers 488 (Baron et al. 1996). This result is expected for long-lived species with low temporal

489 variability in adult survival and an iteroparous reproductive mode that spreads fecundity over 490 several years (Hilde et al. 2020). As a result, stochastic population dynamics are dominated 491 first by a significant deterministic decay term that predicts rapid decline, and second by a 492 strong demographic stochasticity term that accelerates this deterministic decline in small 493 populations. Instead, climate variability is unlikely to slow down or accelerate average 494 stochastic growth. These predictions confirm previous estimates from matrix population 495 models and are consistent with the population decline observed at this isolated site (Ferrière et 496 al. 1996). Climate warming could still benefit the population by improving clutch success, a 497 vital rate positively related to population growth. Yet, average clutch success is already high 498 (around 80%) and our analysis shows that the population growth rate is not as sensitive to 499 components of the life cycle related to reproduction as it is to those related to growth and 500 survival.

501 Our study has implications for assessing the conservation status of the meadow viper 502 and designing measures to protect at least the Mont Ventoux population in France. We show 503 that temporal variability and climatic conditions currently have a limited demographic impact 504 on this population, with warmer temperatures even providing beneficial for breeding. Thus, 505 climate change should not be considered an immediate threat to this population. However, 506 continued monitoring is essential to track the potential effects of more extreme weather 507 conditions, such as summer intense and prolonged hot and dry periods (Madsen et al. 2023). 508 Future assessments should also consider indirect threats posed by climate warming, such as 509 the upward migration of lowland species that could outcompete or prev on meadow vipers, as 510 well as long-term changes in vegetation and habitat. Meanwhile, our analysis underlines the 511 urgent need for strong conservation measures to prevent the expected rapid population of the 512 Mont Ventoux meadow viper. The population's deterministic growth is low, and reversing the 513 decline will require joint improvements of several critical life history traits such as growth and

- 514 survival. Previous research found that growth and late-life survival are enhanced in more
- 515 humid microhabitats (Tully et al. 2020), suggesting that effective conservation actions could
- 516 include habitat restoration. Strict protection measures to decrease disturbance and mortality
- 517 could also be adopted. These could include physical barriers, clear signage, and active
- 518 surveillance to safeguard the population.

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530 Supporting Information

- 531 Supporting Information is available for this article online: Appendix A. Climate variability;
- 532 Appendix B. Model selection tables; Appendix C. Integral projection model

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534 Figure legends

535 Figure 1. Relationship between mean daily maximum temperatures during gestation (red) and 536 during the mating season (green) and the gestation length, calculated as the difference 537 between the parturition date and 10 June, for female meadow vipers at Mont Ventoux, France. 538 Regression lines and confidence bands from the best model in Table B1 are plotted. Note that 539 a cold summer (lowest values of the red data points) is associated with bimodal parturition 540 dates due to reproductive failure (abortions) in several gravid females. 541 542 Figure 2. Relationship between mean daily maximum temperatures during gestation and post-543 partum body condition (residual of a regression between body mass and body size, A) or 544 clutch success (proportion of live newborns in a clutch, B) for female meadow vipers at Mont 545 Ventoux, France. Regression lines and confidence bands from the best models in Tables B3

and B4 are plotted in panels A and B, respectively. In panel B, the size of the data points is

547 proportional to the total clutch size, which ranges from 2 to 8.

548

549 Figure 3. Population viability analysis using an integration projection model (IPM) of the 550 female component of the Mont Ventoux population. A. Elasticity surfaces representing the 551 relative sensitivities of the deterministic growth rate to changes in the kernel function. B. 552 Population viability analysis using a diffusion approximation and simulations of the IBM 553 starting with 100 females randomly drawn from the equilibrium population structure of the 554 deterministic model in the mean environment. Data shown are cumulative quasi-extinction 555 risk during the first 50 years of the simulation for 50,000 population trajectories and three 556 quasi-extinction thresholds (N=1, equivalent to true extinction, N=4 and N=10).

Figure 1



Figure 2





585 **Tables**

586 **Table 1.** Temporal variation in demographic parameters within the study population of the

587 meadow viper at Mont Ventoux, France. Results shown are number of years (N), sample size

588 (*n*), mean (\pm standard deviation for normal distribution and with 95% CI otherwise), best

589 statistical model from a full model including effects of body size, zone and a random year

590 effects, and the relative temporal variation (RTV). RTV measures the proportion of inter-

591 individual variation caused by yearly variation, and was obtained from variance components

592 of the best model.

Demographic parameter	N	п	Mean	Best model	RTV
Reproductive traits				·	
Gestation length*	28	159	90.4 ± 9.81	Year: $\chi^2 = 44.73$, P < 0.001	0.51
Total clutch size	33	220	4.81 ± 1.31	SVL: <i>F</i> _{1,186} = 58.2, P < 0.0001	0.001
				Year: $\chi^2 = 0.001$, P = 0.97	
Clutch success (%)	27	155	0.79 [0.76, 0.82]	SVL: <i>z</i> = -2.01, P = 0.045	0.25
				Year: $\chi^2 = 107.2$, df = 26, P < 0.0001	
Relative clutch mass (g)	24	152	15.4 ± 5.46	Year: $\chi^2 = 5.23$, P = 0.02	0.12
Post-parturition condition (g)	27	157	30.4 ± 9.02	Zone: $F_{1,128} = 14.99$, P < 0.0001	0.27
				Year: $\chi^2 = 12.31$, P < 0.0001	
Offspring traits				•	•
Offspring mass (g)	26	494	2.89 ± 0.45	Mother SVL: $F_{1,114} = 4.69$, P = 0.03	0.11
				Year: $\chi^2 = 188.2$, P < 0.0001	
Offspring condition (residual	26	468	Not applicable (0)	Zone: $F_{1,109} = 3.98$, P = 0.05	0.21
mass)				Year: $\chi^2 = 18.4$, P < 0.0001	
Annual change in body mass and b	ody size				
Body mass change (g)	20	54	12.6 ± 7.29	Zone: $F_{1,32} = 30.2$, P < 0.0001	0.59
				Year: $\chi^2 = 13.8$, P = 0.0002	
Body size growth (mm)	22	57	14.6 ± 11.51	Zone: $F_{1,33} = 10.6$, P = 0.003	0.36
				Year: $\chi^2 = 6.76$, P = 0.009	

593 * Number of days since June 10 (average ovulation and fertilization date)

594 SVL = snout to vent length

596 Supplementary Information

597 Appendix A – Climate variability

598 Temperature data calculated for the three reproductive stages of the meadow vipers at the 599 weather station of the study site were strongly and linearly correlated with temperature data 600 collected at the weather station of Carpentras (Pearson's product moment correlation, n = 21601 years, mating stage: r = 0.95, P < 0.001; gestation stage: r = 0.93, P < 0.001; non-602 reproductive stage: r = 0.91, P < 0.001) and with temperature data calculated with the Safran 603 model for the study site (mating stage: r = 0.97, P < 0.001; gestation stage: r = 0.95, P < 604 0.001; non-reproductive stage: r = 0.94, P < 0.001). To calculate the local weather conditions 605 on the long-term, we thus applied a linear correction factor to the temperature data calculated 606 on the long-term with the Safran model, which was more strongly correlated with local data than data from the Carpentras weather station (mating stage: $F_{1,19} = 274.4$, P < 0.0001, $R^2 =$ 607 0.93; gestation stage: $F_{1,20} = 176.8$, P < 0.0001, $R^2 = 0.90$; non-reproductive stage: $F_{1,20} =$ 608 149.7, P < 0.0001, $R^2 = 0.88$). We obtained similar results when we decomposed data 609 610 calculated for the gestation into three stages (early gestation, mid-gestation, and late gestation: Pearson's product moment correlations, all r > 0.91; linear regressions, all $R^2 > 0.88$). 611 612 Cumulative rainfall data collected at the study site's weather station were also positively 613 correlated with those from Carpentras's weather (Pearson's product moment correlation, n =614 21 years, mating stage: r = 0.77, P < 0.001; gestation stage: r = 0.88, P < 0.001; nonreproductive stage: r = 0.77, P < 0.001) and those calculated with the Safran model (mating 615 616 stages: r = 0.93, P < 0.001; gestation stages: r = 0.94, P < 0.001; non-reproductive stage: r =0.94, P < 0.001). We used a linear correction factor to calculate local rainfall data from data 617 618 calculated with the Safran model, which correlated more strongly with local data than data from the Carpentras weather station (linear regression, mating stage: $F_{1,20} = 126.2$, P < 619

620 0.0001, $R^2 = 0.86$; gestation stage: $F_{1,21} = 152.2$, P < 0.0001, $R^2 = 0.88$; non-reproductive 621 stage: $F_{1,21} = 157.7$, P < 0.0001, $R^2 = 0.89$).

622 Temperature, rainfall, global radiation, and humidity varied markedly among years 623 during the long-term study (Figure A1). Late spring and early summer mean temperatures 624 increased weakly throughout the study period (mating stages: $+0.07 \text{ }^{\circ}\text{C} \pm 0.04 \text{ s.e.}$ per year, 625 $F_{1,33} = 3.16$, P = 0.08; early gestation: + 0.08 °C ± 0.03 s.e. per year, $F_{1,33} = 6.41$, P = 0.02), 626 but this trend wasn't consistent across life stages (other life stages: all P > 0.25, see Figure 627 A1). The rainfall and humidity data showed no consistent temporal trend; instead, they varied 628 in a seemingly random way among years. Global radiation (mean of the cumulative daily 629 global radiation) calculated with the Safran model for the study site increased through time during the study (mating stages: + 15.13 J/cm² ± 4.6 s.e. per year, $F_{1,33} = 10.55$, P = 0.003; 630 gestation: $+7.6 \pm 1.7$ s.e. per year, $F_{1.33} = 20.23$, P < 0.0001, Figure A1). 631

632 Correlations of temperature and rainfall data between reproductive stages within a year 633 were usually small and not significant, but there was a negative correlation between these two 634 environmental factors at the mating and gestation stage, indicating that colder spring and 635 summer climates were also characterized by stronger rainfall (Table A1). Global radiation was 636 consistently positively correlated with temperature and negatively with rainfall within the 637 reproductive stage, and there was also a steady positive correlation between global radiation 638 data across stages within the same year. Humidity was consistently positively correlated with 639 rainfall and temperature within a reproductive stage.

641 Figure A1. We calculated climate conditions for each reproductive year at two life stages of the meadow 642 viper (mating periods [black dots], summer gestation periods [white dots]) from 1980 to 2014. Data 643 from the non-reproductive stage aren't displayed because they had similar temporal trends than data 644 from the gestation stage. (A) Yearly changes in means of maximum daily temperatures (°C); note the 645 exceptionally warm 2003 summer and the warming trend during the mating period represented by the 646 linear regression. (B) Yearly changes in cumulative rainfall (mm); note the wide temporal fluctuations 647 in rainfall during gestation. (C) Yearly changes in daily global radiation (J/cm²) with the increasing trend 648 over time. (D) Yearly changes in mean daily humidity (g/kg).





Table A1. Pairwise correlations of temperature (TP), rainfall (RF), global radiation (GR) and specific humidity (SH) data between reproductive stages calculated

652 each year (upper triangular matrix: Pearson's product moment correlation coefficient, lower triangular matrix: sample size). Reproductive stage: M = mating, G

653 = gestation, and NR = non-reproductive stage of the previous year. Bolded correlations are significant without correction for multiple testing (p < 0.05).

	TP_M	TP_G	TP_NR	RF_M	RF_G	RF_NR	GR_M	GR_G	GR_NR	SH_M	SH_G	SH_NR
TP_M		0.245	-0.055	-0.490	-0.183	-0.051	0.546	0.315	0.170	0.584	0.039	-0.223
TP_G	34		0.056	-0.195	-0.384	-0.019	0.181	0.468	0.087	0.04	0.379	-0.141
TP_NR	34	35		-0.048	-0.011	-0.212	0.104	0.290	0.371	-0.012	-0.139	0.419
RF_M	34	34	34		0.163	0.206	-0.731	-0.168	-0.114	0.192	-0.048	0.112
RF_G	34	35	35	34		0.006	-0.099	-0.446	-0.012	-0.153	0.340	-0.073
RF_NR	34	35	35	34	35		-0.185	-0.274	-0.235	0.134	0.321	0.349
GR_M	34	34	34	34	34	34		0.461	0.403	-0.183	-0.037	-0.053
GR_G	34	35	35	34	35	35	34		0.664	0.173	-0.199	-0.176
GR_NR	34	35	35	34	35	35	34	35		0.038	-0.135	-0.262
SH_M	34	34	34	34	34	34	34	34	34		-0.023	-0.057
SH_G	34	35	35	34	35	35	34	35	35	35		0.141
SH_NR	34	35	35	34	35	35	34	35	35	35	35	

657 Appendix B – Model selection tables

- 658 Notations in Appendix B are k (number of model parameters), LogLikelihood (natural logarithm of the likelihood function), AICc (Akaike
- 659 Information Criterion corrected for small sample size), Δ_i (difference between the AIC value of the best model and the AIC value for each other
- models), w_i (Akaike weight), ER (evidence ratio), R²m (marginal pseudo-R2 for the mixed-effects model) and R²c (conditional pseudo-R2 for the
- 661 mixed-effects model). Abbreviations for the descriptions of the covariate are the same as in Table A1. First part corresponds to the kind of
- 662 climate variable: TP = temperature, GR = global radiation, and SH = specific humidity. Second part corresponds to the kind of life stage: M =
- 663 mating, G = gestation, and NR = non-reproductive stage of the previous year.

Table B1. Full set of linear models examining effect of climate variables on gestation length in the meadow viper population at Mont Ventoux,

666	France. The best set of models	(models whose cumulative	Akaike weight ≤ 0.95) is in bold.	Yellow filling indicates models with $\Delta_i < 2$.
			$U = \prime$	

Model rank	Model name	k	-LogLikelihood	AICc	Δ_{i}	Wi	ER	R ² m	R ² c
1	TP_M+TP_G	5	-557.53	1125.46	0.00	0.47		0.27	0.47
2	TP_NR+TP_M+TP_G	6	-556.58	1125.71	0.25	0.41	1.13	0.28	0.46
3	TP_M	4	-561.17	1130.60	5.14	0.04	13.06	0.19	0.48
4	TP_NR+TP_G	5	-560.28	1130.94	5.48	0.03	15.51	0.17	0.48
5	TP_G	4	-561.61	1131.49	6.03	0.02	20.34	0.15	0.50
6	TP_NR+TP_M	5	-560.91	1132.21	6.75	0.02	29.15	0.19	0.48
7	GR_M	4	-562.57	1133.40	7.93	0.01	52.81	0.12	0.49
8	1	3	-565.89	1137.94	12.48	0.00	512.75	0.00	0.51
9	TP_NR	4	-565.37	1138.99	13.53	0.00	868.01	0.03	0.51
10	GR_G	4	-565.37	1139.00	13.54	0.00	872.67	0.02	0.51
11	SH_M	4	-565.62	1139.50	14.04	0.00	1117.98	0.01	0.50
12	GR_NR	4	-565.65	1139.56	14.09	0.00	1149.60	0.01	0.51
13	SH_NR	4	-565.80	1139.87	14.41	0.00	1343.79	0.00	0.51
14	SH_G	4	-565.87	1140.00	14.54	0.00	1437.36	0.00	0.51
15	SH_NR+SH_M	5	-565.56	1141.51	16.05	0.00	3055.19	0.01	0.50
16	SH_M+SH_G	5	-565.61	1141.61	16.14	0.00	3203.81	0.01	0.50
17	SH_NR+SH_G	5	-565.79	1141.97	16.51	0.00	3839.25	0.00	0.51
18	SH_NR+SH_M+SH_G	6	-565.55	1143.65	18.19	0.00	8906.16	0.01	0.50

Table B2. Full set of linear models examining effect of climate variables on relative clutch mass in the meadow viper population at Mont

669	Ventoux, France.	The best set of models	(models whose	cumulative Aka	ike weight \leq	0.95) is in bold.	Yellow filling	indicates models	with $\Delta_i < 2$.
	,				<i>u</i> =	/	<i>u</i>		

Model rank	Model name	k	-LogLikelihood	AICc	Δ_{i}	Wi	ER	R2m	R2c
1	GR_NR	5	-453.22	916.85	0.00	0.25		0.21	0.29
2	GR_G	5	-453.80	918.01	1.16	0.14	1.79	0.19	0.30
3	1	4	-455.13	918.54	1.69	0.11	2.33	0.17	0.29
4	SH_G	5	-454.08	918.57	1.73	0.10	2.37	0.19	0.30
5	SH_M+SH_G	6	-453.84	920.26	3.41	0.05	5.50	0.19	0.30
6	TP_M	5	-454.95	920.32	3.47	0.04	5.67	0.17	0.29
7	SH_M	5	-455.05	920.51	3.66	0.04	6.23	0.16	0.29
8	GR_M	5	-455.05	920.51	3.67	0.04	6.26	0.17	0.29
9	SH_NR+SH_G	6	-453.99	920.56	3.71	0.04	6.41	0.19	0.30
10	SH_NR	5	-455.09	920.58	3.74	0.04	6.47	0.17	0.29
11	TP_G	5	-455.10	920.62	3.77	0.04	6.59	0.16	0.29
12	TP_NR	5	-455.12	920.65	3.80	0.04	6.68	0.16	0.29
13	SH_NR+SH_M+SH_G	7	-453.69	922.16	5.31	0.02	14.22	0.20	0.30
14	TP_M+TP_G	6	-454.86	922.30	5.45	0.02	15.29	0.16	0.29
15	TP_NR+TP_M	6	-454.95	922.47	5.63	0.01	16.68	0.17	0.29
16	SH_NR+SH_M	6	-454.98	922.54	5.69	0.01	17.22	0.17	0.29
17	TP_NR+TP_G	6	-455.09	922.77	5.92	0.01	19.33	0.16	0.29
18	TP_NR+TP_M+TP_G	7	-454.86	924.50	7.65	0.01	45.90	0.16	0.29

Table B3. Full set of linear models examining effect of climate variables on post-parturition condition in the meadow viper population at Mont

673	Ventoux, France.	The best set of models	(models whose	cumulative Akaike	e weight ≤ 0.95	5) is in bold.	Yellow filling indicates	models with $\Delta_i < 2$.
	· · · · · · · · · · · · · · · · · · ·				\mathcal{U} =	/	0	

Model rank	Model name	k	-LogLikelihood	AICc	Δ_{i}	Wi	ER	R2m	R2c
1	TP_NR+TP_G	7	92.11	-169.45	0.00	0.18		0.81	0.84
2	TP_NR	6	90.76	-168.94	0.51	0.14	1.29	0.80	0.84
3	TP_NR+TP_M+TP_G	8	92.73	-168.45	1.01	0.11	1.65	0.82	0.84
4	GR_NR	6	90.42	-168.27	1.19	0.10	1.81	0.80	0.83
5	TP_G	6	90.42	-168.25	1.20	0.10	1.82	0.81	0.84
6	TP_M+TP_G	7	91.43	-168.08	1.37	0.09	1.99	0.81	0.84
7	1	5	88.74	-167.06	2.39	0.05	3.30	0.79	0.84
8	TP_NR+TP_M	7	90.90	-167.03	2.43	0.05	3.36	0.81	0.84
9	GR_G	6	89.61	-166.65	2.80	0.04	4.06	0.80	0.83
10	TP_M	6	89.06	-165.53	3.92	0.03	7.09	0.80	0.84
11	SH_M	6	89.00	-165.43	4.02	0.02	7.48	0.80	0.84
12	GR_M	6	88.88	-165.18	4.27	0.02	8.46	0.79	0.84
13	SH_NR	6	88.75	-164.91	4.54	0.02	9.67	0.79	0.84
14	SH_G	6	88.74	-164.91	4.54	0.02	9.69	0.79	0.84
15	SH_NR+SH_M	7	89.00	-163.23	6.22	0.01	22.43	0.80	0.84
16	SH_M+SH_G	7	89.00	-163.23	6.22	0.01	22.44	0.80	0.84
17	SH_NR+SH_G	7	88.75	-162.73	6.72	0.01	28.81	0.79	0.84
18	SH_NR+SH_M+SH_G	8	89.00	-161.00	8.45	0.00	68.39	0.80	0.84

- **Table B4.** Full set of binomial models examining effect of climate variables on clutch success in the meadow viper population at Mont Ventoux,
- France. The best set of models (models whose cumulative Akaike weight ≤ 0.95) is in bold. Yellow filling indicates models with $\Delta_i < 2$. The set

Model rank	Model name	k	-LogLikelihood	AICc	Δ_{i}	Wi	ER	R2m	R2c
1	TP_NR+TP_G	5	-157.85	326.10	0.00	0.21		0.06	0.10
2	TP_G	4	-159.13	326.53	0.43	0.17	1.24	0.04	0.09
3	SH_NR	4	-159.76	327.79	1.69	0.09	2.32	0.02	0.09
4	TP_NR+TP_M+TP_G	6	-157.85	328.27	2.16	0.07	2.95	0.06	0.10
5	TP_M+TP_G	5	-159.08	328.56	2.46	0.06	3.42	0.05	0.09
6	1	3	-161.27	328.70	2.60	0.06	3.66	0.00	0.07
7	SH_NR+SH_M	5	-159.32	329.04	2.94	0.05	4.35	0.03	0.09
8	SH_M	4	-160.46	329.19	3.09	0.05	4.69	0.01	0.08
9	SH_NR+SH_G	5	-159.50	329.40	3.30	0.04	5.20	0.03	0.09
10	TP_NR	4	-160.67	329.60	3.50	0.04	5.76	0.09	0.08
11	TP_M	4	-160.97	330.21	4.11	0.03	7.81	0.10	0.08
12	GR_NR	4	-161.03	330.33	4.23	0.03	8.27	0.01	0.08
13	SH_G	4	-161.09	330.45	4.34	0.02	8.78	0.01	0.07
14	SH_NR+SH_M+SH_G	6	-158.94	330.45	4.35	0.02	8.78	0.04	0.09
15	GR_G	4	-161.12	330.50	4.40	0.02	9.02	0.01	0.07
16	SH_M+SH_G	5	-160.14	330.68	4.58	0.02	9.89	0.02	0.07
17	TP_NR+TP_M	5	-160.47	331.35	5.25	0.02	13.82	0.01	0.09

678 of models did not include GR_M effect due to lack of convergence.

Table B5. Full set of linear models examining effect of climate variables on offspring mass at birth in the meadow viper population at Mont

683	Ventoux, France.	The best set of models (n	nodels whose cumulative	Akaike weight ≤ 0.95) is in bold.	Yellow filling indicates models	s with $\Delta_i < 2$.
	· · · · · · · · · · · · · · · · · · ·			$\mathcal{U} = \mathcal{V}$	0	

Model rank	Model name	k	-LogLikelihood	AICc	Δ_{i}	Wi	ER	R2m	R2c
1	1	6	-192.89	397.95	0.00	0.15		0.05	0.61
2	GR_NR	7	-192.30	398.84	0.88	0.09	1.55	0.06	0.61
3	TP_NR	7	-192.33	398.89	0.94	0.09	1.60	0.06	0.61
4	TP_G	7	-192.41	399.06	1.10	0.08	1.73	0.05	0.61
5	GR_M	7	-192.59	399.41	1.46	0.07	2.07	0.05	0.60
6	TP_NR+TP_G	8	-191.59	399.48	1.52	0.07	2.14	0.07	0.61
7	TP_M	7	-192.68	399.58	1.63	0.07	2.25	0.05	0.60
8	GR_G	7	-192.83	399.89	1.94	0.06	2.64	0.05	0.61
9	SH_M	7	-192.87	399.97	2.02	0.05	2.74	0.05	0.61
10	SH_NR	7	-192.88	399.99	2.03	0.05	2.76	0.05	0.61
11	SH_G	7	-192.89	400.01	2.06	0.05	2.80	0.05	0.61
12	TP_NR+TP_M	8	-192.17	400.64	2.69	0.04	3.83	0.06	0.61
13	TP_M+TP_G	8	-192.31	400.91	2.95	0.03	4.38	0.05	0.60
14	TP_NR+TP_M+TP_G	9	-191.55	401.47	3.51	0.03	5.79	0.07	0.61
15	SH_NR+SH_M	8	-192.86	402.01	4.05	0.02	7.58	0.05	0.61
16	SH_M+SH_G	8	-192.87	402.04	4.08	0.02	7.69	0.05	0.61
17	SH_NR+SH_G	8	-192.88	402.05	4.10	0.02	7.76	0.05	0.61
18	SH_NR+SH_M+SH_G	9	-192.85	404.08	6.12	0.01	21.37	0.05	0.61

Table B6. Full set of linear models examining effect of climate variables on offspring body condition at birth in the meadow viper population at

687	Mont Ventoux,	France. The b	est set of models	(models whos	se cumulative A	Akaike weight < ((0.95) is in bold.
				(

Model rank	Model name	k	-LogLikelihood	AICc	Δ_{i}	Wi	ER	R2m	R2c
1	GR_G	8	10.24	-4.17	0.00	0.54		0.63	0.80
2	TP_G	8	8.72	-1.13	3.04	0.12	4.58	0.63	0.80
3	GR_NR	8	8.18	-0.04	4.13	0.07	7.87	0.61	0.80
4	TP_M+TP_G	9	8.92	0.56	4.73	0.05	10.63	0.63	0.80
5	TP_NR+TP_G	9	8.82	0.75	4.92	0.05	11.70	0.63	0.80
6	1	7	6.38	1.48	5.66	0.03	16.91	0.61	0.81
7	TP_NR+TP_M+TP_G	10	9.08	2.32	6.49	0.02	25.68	0.63	0.80
8	TP_M	8	6.94	2.43	6.60	0.02	27.17	0.62	0.81
9	GR_M	8	6.88	2.55	6.72	0.02	28.75	0.62	0.81
10	SH_G	8	6.70	2.92	7.09	0.02	34.69	0.62	0.81
11	TP_NR	8	6.62	3.06	7.24	0.01	37.26	0.61	0.80
12	SH_M	8	6.41	3.50	7.67	0.01	46.32	0.61	0.81
13	SH_NR	8	6.38	3.55	7.72	0.01	47.52	0.61	0.81
14	TP_NR+TP_M	9	7.32	3.76	7.93	0.01	52.76	0.61	0.80
15	SH_M+SH_G	9	6.71	4.98	9.15	0.01	96.97	0.62	0.81
16	SH_NR+SH_G	9	6.70	5.00	9.17	0.01	98.07	0.62	0.81
17	SH_NR+SH_M	9	6.41	5.57	9.74	0.00	130.45	0.61	0.81
18	SH_NR+SH_M+SH_G	10	6.71	7.07	11.24	0.00	275.50	0.62	0.81

689 Appendix C – Integral projection model

690 Model rationale

The integral projection model (IPM) assumes that the life history is structured by body size (here, snout-vent length), hereafter called *y*, such that snakes differ by *y* only and *y* is the sole determinant of their vital rates. The population can then be described by a function N(y,t) where N(y,t)dy is the discrete number of individuals between size values *y* and y + dy at time *t*. The total, finite population size is called N(t). To describe this population in a fluctuating environment, we used a stochastic IPM formulated by Vindenes et al. (2011):

697
$$N(y,t+1) = \int_{\Omega} K(y,x,Z_t) N(x,t) dx,$$
 (C1)

698
$$K(y, x, Z_t) = s(x, Z_t) f_s(y, x, Z_t) + b(x, Z_t) f_b(y, x, Z_t)$$

699 where *K* is a stochastic projection kernel, Z_t is a random vector describing parameter values at 700 time *t* and thus the environmental state, and Ω is the domain of possible values for body size (Rees 701 and Ellner 2009).

702 The kernel describes transition rates from state x at time t to state y at time t+1 in environment Z. According to equation (C1), the kernel is decomposed into (1) the survival-growth 703 kernel where $s(xZ_t)$ is the survival probability of an individual with trait value x, and 704 $f_s(y, x, Z_t)dy$ is the probability of reaching trait value between y and y + dy at time t+1 for an 705 individual of trait value x, and (2) the fecundity kernel where $b(x, Z_t)$ is the fecundity of an 706 individual with trait value x, and $f_b(y, x, Z_t)$ is the probability density function of the trait value of 707 708 offspring. Thus, the stochastic model described by equation (C1) is similar the deterministic model 709 of Easterling et al. (2000) conditional on Z.

710 This IPM can be considered as a stochastic matrix projection model (MPM) with an infinite number of discrete classes. According to the seminal paper by Easterling et al. (2000), most of the 711 712 properties of MPMs can therefore be generalised to IPMs, including the calculation of the 713 deterministic population growth rate λ , equilibrium population structure, reproductive values and 714 deterministic elasticities of λ . Here, we used a numerical method to simulate the deterministic IPMs by discretising the state-space Ω into C classes of the same width Δy and computing 715 integrals using Simpson's 3/8 method, a more accurate numerical integration method than the 716 717 standard mid-point rule (Merow et al. 2014).

For stochastic environments and finite populations, we used individual based simulations and considered that Z_t is a vector of year-specific parameters and assumed that parameter values vary randomly over time according to the random effects model of Rees and Ellner (2009). This implies that elements of the stochastic kernel of the IPM are drawn randomly each year from some parametric statistical distributions. The dynamics of the expected population size at time *t*+1 given population size at time *t* can then be written as:

724
$$E[N(t+1)|N(t)] = \int_{\Omega} \int_{\Omega} \overline{k}(y,x)N(x,t)dxdy$$
(C2)

725 where $\overline{k}(y, x)$ is the mean kernel defined by averaging the stochastic kernel over all possible 726 environmental state values. This dynamics is characterised by an expected growth rate $\overline{\lambda}$ and a 727 stable size structure $\overline{u}(x)$.

An important property of this stochastic IPM is the long-run logarithmic growth rate, denoted $\ln \lambda_s$, which describes the asymptotic exponential growth rate of the population size after a sufficiently long time (Lande et al. 2003; Rees and Ellner 2009; Tuljapurkar 1990). The long-run logarithmic growth rate can be approximated assuming a small environmental noise, which implies

that the population stays close to its stable distribution (e.g., for IPM Rees and Ellner 2009).
Vindenes et al. (2011) showed that the first-order approximation of the long-run growth rate of the
stochastic IPM writes like:

735
$$\ln \lambda_s \approx \ln \overline{\lambda} - \frac{\sigma_e^2}{2\overline{\lambda}^2} - \frac{\sigma_d^2}{2\overline{\lambda}^2 V}, \qquad (C3)$$

where $\overline{\lambda}$ is the deterministic population growth rate of the mean kernel, σ_e^2 is the environmental 736 variance (i.e., a consequence of between-year deviation from the mean kernel of the average 737 individual contribution to total reproductive value), σ_d^2 is the demographic variance (i.e., a 738 consequence within-year deviation from the mean of the year of the individual contribution to total 739 740 reproductive value) and V is the total reproductive value (i.e., equivalent to population size when the population reaches the stable distribution). We calculated σ_e^2 by numeric simulations of the 741 growth rate, estimated σ_d^2 from the slope of the regression of the mean exponential growth rate 742 against the inverse population size, and then simulated model (C2) with a diffusion approximation 743 744 of long-run growth rate following the numerical methods of Vindenes et al. (2011).

745 Model parameterization

746 Our female life cycle for the meadow viper counts individuals from the age of one year and assumes annual reproduction followed by annual survival and body growth of juveniles, and the 747 annual survival, body growth and sexual maturation of immature individuals (until six years old, 748 749 equivalent to 7 age classes of immature individuals). Juvenile survival increases with offspring 750 mass at birth, which itself depends on offspring body size (snout-vent length, mm) that is influenced 751 by maternal body size and total clutch size. Maturation probability depends on both body size and age of the female, and occurs around 4 to 6 years old. Once maturity is reached, females alternate 752 753 regularly breeding and non-breeding years (equivalent to two breeding classes of mature

individuals) and their fecundity increases with body size. To calculate the number of viable 754 offspring produced by breeding females each year, we estimated fecundity (total clutch size), clutch 755 756 success (percentage of viable offspring within a clutch) and female sex ratio. This complex life cycle was modelled with an IPM including both a continuous size structure and nine discrete 757 categories. The first seven discrete categories were age classes (from 0 to 6 years) for immature 758 759 individuals, and the two last discrete categories included the breeding and non-breeding states for mature individuals. There is significant environmental variation for annual juvenile survival, body 760 size at birth and reproduction through clutch success. This life history is based on earlier results 761 (Baron et al., 2010a, Baron et al., 2013, Baron et al., 2010b) and results from the current study. 762

The IPM for this complex life cycle can be written like:

764
$$n(y,t+1) = \int_{Y} k(y,x) \times n(x,t) \times dx$$
, (C4)

where n(y,t) is a vector of functions describing the size distribution in each class, and k is a matrix of functions whose elements are kernel elements of a classical IPM for each transition between classes. More precisely, we can write down the matrix k like:

$$768 \qquad k(y,x) = \begin{pmatrix} 0 & 0 & 0 & 0 & 0 & 0 & 0 & k_F & 0 \\ k_{0\to1} & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & k_{1\to2} & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & k_{2\to3} & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & k_{3\to4} & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & k_{4\to5} & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & k_{5\to6} & 0 & 0 & 0 \\ k_{0\toB} & k_{1\toB} & k_{2\toB} & k_{3\toB} & k_{4\toB} & k_{5\toB} & k_{B\toB} & k_{NB\toB} \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & k_{B\toNB} & k_{NB\toNB} \end{pmatrix}$$
(C5)

769 where elements of this matrix are listed below:

770
$$k_{i \to j}(y, x) = (1 - \prod_{m} (y, i+1)) \times f_{Im}(y, x) \times \phi_{Im}(y, x)$$

771
$$k_F(y,x) = \left[\sum_{k=1}^{F_{max}} \Pi(F_0(x) = k) \times k \times f_F(y,x \mid F_0(x) = k)\right] \times p_{cs}(x) \times p_f \times \phi_0(m(y))$$

772
$$k_{i\to B}(y,x) = \prod_{m} (y,i+1) \times f_i(y,x) \times \phi_{lm}$$

773
$$k_{B\to B}(y,x) = P_{B\to B} \times f_B(y,x) \times \phi_A$$

774
$$k_{NB \to B}(y, x) = P_{NB \to B} \times f_{NB}(y, x) \times \phi_{A}$$

775
$$k_{B \to NB}(y, x) = P_{B \to NB} \times f_B(y, x) \times \phi_A$$

776
$$k_{NB \to NB}(y, x) = P_{NB \to NB} \times f_{NB}(y, x) \times \phi_A$$

777

Parameters values for the different terms of these elements are provided in Table C1 averaging across the two study sites. The stochastic model included inter-annual variation in clutch success, size at birth and size growth of non-reproductive females (this study) as well as early juvenile survival (Baron et al., 2010a, Baron et al., 2013, Baron et al., 2010b). The stochastic version was analysed with an individual based model in Python whereas the deterministic version was fitted with Matlab. Default parameter values of the deterministic model provided a good fit to the average size distribution in the study population (see Figure C1)

785

Figure C1. Comparison between the size distribution predicted by the IPM and the observed size distribution in the pooled sample of females corrected for unequal capture probabilities. The two statistical distributions looked similar although the IPM tended to predict more of the small juvenile size classes and less of the old adults' size classes than observed (χ^2 test, P < 0.0001).

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Table C1: Model parameters when size is estimated by snout-vent length (SVL, mm) for *Vipera ursinii* ursinii at Mont Ventoux, France. The table provides the mean and error of the mean in the scale of the statistical model used to estimate parameters, as well as inter-annual SD, statistical procedure / software and kinds of distribution (glm: generalized linear model, lm: linear model, glmmPQL: generalized linear mixed model with penalized quasi-likeihood, vglm: vector generalized linear model, lme: liner mixed model,

800 MARK CJS: Cormak-Jolly-Seber open population model including multi-state models).

Trait	Scale	Estimate	SE	Inter-annual SD	Procedure	Distribution		
Sexual maturation probability - $\prod_{n}(y,i)$								
Intercept	cept Logit		5.88	0	glm	Binomial		
Slope age <i>i</i> Logit		2.072	0.530	0				
Slope SVL y	Logit	0.050	0.017	0				
Size of immature femal	es at time t	$+1 - f_{Im}(y, x)$.)	•				
Intercept	Identity	83.646	10.82	0	lm	Gaussian		
Slope SVL <i>x</i>	De SVL x Identity		0.043	0				
Residual variance*	Power	321097	NA					
Annual survival of imm	ature femal	es - ϕ_{Im}		•				
Mean value	Logit	1.202	0.467	0	MARK CJS	Binomial		
Fecundity (total clutch	size) - $F_0(x)$)	•					
λ - Mean value	elogit	-1.665	0.294	0	vglm	Generalised		
Θ - Intercept	log	-1.090	0.285	0		Poisson		
Θ - Slope SVL <i>x</i>	log	0.007	0.0007	0				
Clutch success - $p_{cs}(x)$)							
Intercept	Logit	3.662	1.822	1.139	glmmPQL	Binomial		
Slope maternal SVL <i>x</i>	Logit	-0.024	0.323	0				
Residual variance	Logit	1.067	NA					
Size at birth given fecu	ndity - $f_F(y)$	$V, x \mid F_0(x))$		•				
Intercept	Identity	116.895	4.14	4.064	lme	Gaussian		
Slope maternal SVL <i>x</i>	lope maternal SVL x Identity		0.012	0				
Slope fecundity $F_0(x)$	Identity	-1.389	0.351	0				
Residual variance	Identity	6.143	NA					
Female sex ratio - p_f			1	1				
Intercept	Logit	-0.044	0.1054		glm	Binomial		

Juvenile survival as a fu	nction of st	andardised bo	ody mass	at birth - $\phi_0(m(y))$		
Intercept	Logit	-0.442	0.505	0.269	MARK CJS	Binomial
Slope body mass $m(y)$	Logit	0.837	0.539	0		
Relationship between ma	ass at birth	and size at bi	rth - $m(y)$			
Intercept	Identity	-3.393	0.276	0	lm	Gaussian
Slope size at birth y	Identity	0.047	0.002	0		
Residual variance	Identity	0.2933	NA			
Transition from breedin	g to non-br	eeding state -	$P_{B \rightarrow NB}$	1	1	1
Mean value	Logit	3.200	0.524	0	MARK CJS	Binomial
Size of breeding females	at time t +	$1 - f_B(y, x)$	1	1	1	
Intercept	Identity	21.559	14.76	0	lm	Gaussian
Slope SVL <i>x</i>	Identity	0.9514	0.039	0		
Residual variance	Identity	9.04	NA			
Adult survival - ϕ_A	I		1	-	-	
Mean value	Logit	0.789	0.116	0	MARK CJS	Binomial
Transition from non-bre	eding to br	eeding state -	$P_{NB \rightarrow B}$	-	-	
Mean value	Logit	1.756	0.391	0	MARK CJS	Binomial
Size of non-breeding fen	nales at tim	e $t+1 - f_{NB}$	(y, x)	-	-	
Intercept	Identity	84.821	21.88	1.23	lm	Gaussian
Slope SVL <i>x</i>	Identity	0.8204	0.058	0		
Residual variance	Identity	11.43	NA			

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* Residual variance is modelled like a power function of predicted values (power exponent: -1.79)