1	Colour-assortative mating in a colour-polymorphic lacertid lizard				
2	¹ Pérez i de Lanuza, Guillem; ¹ Font, Enrique and ^{1, 2} Carazo, Pau				
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4	¹ Cavanilles Institute of Biodiversity and Evolutionary Biology, University of Valencia				
5	² Edward Grey Institute, Department of Zoology, University of Oxford				
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7	Short running title: Colour assortative mating in Podarcis muralis				
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9	Correspondence: G. Pérez i de Lanuza, Ethology Lab, Institut Cavanilles de				
10	Biodiversitat i Biologia Evolutiva, University of Valencia, APDO. 22085, 46071				
11	Valencia, Spain.				
12	Tel.: +34 963543659; fax: +34 963543670				

13 e-mail: guillem.perez-lanuza@uv.es

14 Abstract

15 Colour polymorphisms are common in lizards, which provide an excellent model system to study their evolution and adaptive function. The lacertid genus Podarcis is 16 particularly interesting because it comprises several polymorphic species. Previous 17 18 studies with lacertid lizards have tried to explain the maintenance of colour polymorphisms by correlational selection between colour morphs and several 19 phenotypic traits. Particular attention has been paid to their putative role as signals 20 21 reflecting alternative reproductive strategies under frequency-dependent selection, but 22 the relationship between mating patterns and colour polymorphism has not been previously considered. In this study, we use longitudinal behavioural data obtained 23 24 during six consecutive breeding seasons (2006-2011) in a free-ranging polymorphic population of *Podarcis muralis* lizards to examine the hypothesis that lizards mate 25 assortatively by colour. We provide spectrophotometric data that confirm the existence 26 27 of discrete colour morphs and show that morphs are ontogenetically stable once they develop fully in sexually mature individuals. We also present data on the year-to-year 28 29 variation of relative morph frequencies. Finally, we provide evidence that, over a six-30 year period, homomorphic male-female pairs in the wild were significantly more common than heteromorphic pairs. Taken together, our results suggest that colour 31 32 assortative mating may be involved in the maintenance of discrete colour morphs in this and other lacertid species. 33

Keywords: alternative reproductive strategies; assortative mating; colour variation;
polymorphism

36 Introduction

37	A major challenge in evolutionary biology is to understand the genetic mechanisms
38	and evolutionary processes involved in the maintenance of phenotypic variation,
39	including colour polymorphisms (Roulin 2004; Chunco et al. 2007; Pryke and Griffith
40	2007; McKinnon and Pierotti 2010). Studies with lizards have contributed importantly
41	to this endeavour, and suggest that the lizard model is an ideal one in which to test
42	hypotheses and predictions about the evolution and maintenance of colour
43	polymorphisms (Sinervo and Lively 1996; Zamudio and Sinervo 2000; Vercken et al.
44	2007). Colour polymorphisms are common in lacertids (Arnold and Oveden 2002),
45	and typically consist of lizards exhibiting either white or orange ventral colourations
46	(i.e. throat and/or belly), although some populations feature an additional yellow
47	morph as well as several rare intermediate phenotypes. Polymorphic populations of
48	lizards seem to be particularly abundant in the lacertid genus Podarcis (P.
49	melisellensis, Huyghe et al. 2007, 2009a,b, 2010a,b; P. muralis, Cheylan 1988; Sacchi
50	et al. 2007a,b, 2009; Calsbeek et al. 2010; Galeotti et al. 2010; Font et al. 2010; P.
51	gaigeae, Runemark et al. 2010; P. vaucheri and P. liolepis, G. Pérez i de Lanuza,
52	unpublished data), which makes it an excellent system in which to study the evolution,
53	maintenance, and adaptive function of genetic colour polymorphisms.
54	While different evolutionary processes may underlie colour polymorphisms

(Roulin 2004; Bond 2007; Gray and McKinnon 2007; Roulin and Bize, 2007), work
on lizards has mainly focused on the search for phenotypic syndromes associated with

- 57 discrete colour polymorphisms (e.g., Sinervo and Calsbeek 2006; Calsbeek et al.
- 58 2010). Several studies have explored potential differences across colour morphs in a
- 59 number of life-history, behavioural, performance, morphological, ecological and

60 physiological traits. In particular, much attention has been paid to the possibility that 61 lacertid colour polymorphisms reflect the existence of alternative reproductive strategies that would be maintained by frequency-dependent selection (Sinervo and 62 Lively 1996; Sinervo et al. 2007). In P. melisellensis, males of different colour morphs 63 have been found to differ in several phenotypic traits (i.e. morphology, fighting 64 ability, corticosterone levels, haemoparasite infection levels, immune response) that 65 66 hint at the existence of alternative reproductive strategies (Huyghe et al. 2007, 2009a, 2010b). In the wall lizard (*P. muralis*) male morphs do not differ in aggressive levels 67 or fighting success, although immune response and susceptibility to stress in captivity 68 69 appear to covary with colour morphs (Sacchi et al. 2007b, 2009; Galeotti et al. 2010). Calsbeek et al. (2010) recently reported differences among P. muralis morphs in 70 71 haemoparasite infection rate, intensity, and probability of survival. These authors 72 proposed that *P. muralis* morphs experience different multivariate selection pressures, and suggested that such correlational selection may have favoured the evolution of 73 74 alternative optimal morph-specific phenotypes (Lande and Arnold 1983; Forsman et al. 2008). However, the fact that multiple phenotypic optima may underlie alternative 75 behavioural strategies does not explain why each strategy should be associated with a 76 77 different ventral colouration (Roulin 2004); indeed, there is no experimental confirmation of the presumed signalling role of ventral colourations or of the 78 79 information they make available to potential receivers. In conclusion, we are still far from identifying the genetic and evolutionary processes underlying the maintenance of 80 colour polymorphisms in lacertid lizards. 81

Mate choice has been put forward as one of the evolutionary mechanisms that can contribute to maintaining colour polymorphisms (Roulin 2004; Pryke and Griffith 2007; Puebla et al. 2007; Reynolds and Fitzpatrick 2007; Elmer et al. 2009). Although

the evolution of monotonic directional preferences is predicted for condition-85 86 dependent colour traits, in polymorphic systems colour-based mate choice decisions may be morph-specific. This will happen when the reproductive fitness of mating with 87 a given morph is dependent on the morph of both sexes (Roulin 2004; Pryke and 88 Griffith 2009), and will be reflected by the existence of nonrandom assortative or 89 disassortative mating between different colour morphs (Lank 2002; Galeotti et al. 90 91 2003; Roulin 2004; Gray and McKinnon 2007; Pryke and Griffith 2007; Roulin and Bize 2007). 92

93 Here we address the possibility that mate choice patterns (i.e. mate recognition/assessment sensu Johansson and Jones 2007) may be non-random with 94 95 respect to colour morph. To this end, we used data collected over six consecutive breeding seasons in a free-ranging polymorphic population of *P. muralis*: 1) to 96 objectively evaluate (i.e. using spectrophotometric methods) the existence of discrete 97 colour morphs, 2) to assess the ontogenetic stability of colour morphs, 3) to study 98 inter-seasonal variation in relative morph frequencies, and 4) to analyse the existence 99 100 of assortative colour-based mating between different morphs.

101 Materials and methods

102 *Study population*

103 We studied a white-yellow-orange polymorphic population of *P. muralis* from the

104 Cerdanya valley in the south-eastern Pyrenees. In this population, males and females

typically exhibit three alternative pure-colour morphs (white, W; yellow, Y; orange,

106 O), although some individuals show intermediate phenotypes consisting of a mosaic of

107 white and orange scales (WO) or a mosaic of yellow and orange scales (YO). Despite

extensive sampling, we did not find lizards with an intermediate white and yellow 108 phenotype as described by Calsbeek et al. (2010). Whereas in males the ventral 109 colourations extend over the throat and belly, in females yellow and orange 110 colourations are restricted to the throat (i.e. the belly is always white). All the lizards 111 112 included in this study came from a homogeneous, continuous population, so our estimates of morph frequency are not confounded by inter-population differences in 113 morph frequencies, which seem common in this species (Cheylan 1988; Sacchi et al. 114 2007a; G. Pérez i de Lanuza, unpublished results). 115

116 Colour morph characterization, colour morph development and polymorphism117 stability

Our main study area was a 2.1 Ha continuous patch of abandoned cultivated terraces 118 with artificial stonewalls and sparse vegetation mainly consisting of ashes, rosebushes, 119 hawthorns, and blackthorns. Each spring between 2006 and 2011, we performed field 120 surveys in this area during 4-5 weeks from late May to early July. Overall, we 121 captured 770 animals for which we determined colour morph (eye-based assignment), 122 sex, and age (i.e. juvenile, subadult or adult). We measured body size (as snout-vent 123 124 length, SVL), body mass, and head width (HW) in all the lizards, and throat colouration in a subsample of adult males and females. To obtain objective 125 measurements of throat colouration we used a USB2000 portable spectrometer with a 126 PX-2 Xenon strobe lamp (Ocean Optics Inc., Dunedin, FL, USA) and standard 127 spectrophotometric techniques (for details see Font et al. 2009; Pérez i de Lanuza and 128 Font 2011). We restricted colour analyses to the 300-700 nm range, which 129 encompasses the visual spectrum of diurnal lizards (Fleishman et al. 1993, 1997; 130 Loew et al. 2002). We measured brightness (Q) and hue (H) according to Endler's 131

Segment Classification method (1990). Additionally, we calculated medium 132 wavelength chroma (MC) as R400-600/R300-700, where R400-600 and R300-700 are 133 the sums of the per cent reflectance between 400 and 600 nm and between 300 and 134 700 nm, respectively. We chose this chroma variable because chromatic differences 135 among colour morphs are due mainly to variation in reflectance between 400 and 600 136 nm (see Figure 1). We also calculated the spectral location of the median reflectance 137 $(R_{50}, i.e.$ the difference between maximum and minimum reflectance divided by two; 138 Marshall et al. 2003). 139

Spectral data were analysed by fitting a two-way ANOVA model with morph 140 and sex as fixed factors. In those cases in which we could not assume data to be 141 142 normally distributed we used the Scheirer-Ray-Hare test (Sokal and Rohlf 2009). Finally, we used one-way ANOVA or Kruskal-Wallis (i.e. when heteroscedasticity 143 and/or normality could not be safely assumed) tests to look for inter-morph 144 morphometric differences in SVL and body condition. Body condition (BCI) was 145 calculated as the body mass residuals after regressing body mass against SVL (Green 146 147 2001). We restricted this analysis to adult lizards with fully developed colouration (i.e. SVL > 56 mm; see below). 148

Before releasing lizards back at their capture locations, all the individuals were marked by toe-clipping for subsequent identification. We used toe-clipping because this is the most adequate and ethically sound method for durable marking in lizards of this size range (for full details see Perry et al. 2011). Briefly, we clipped toes by cutting their distal two-thirds with a pair of sharp surgical scissors. We clipped a maximum of two toes and always tried to select small digits that did not usually draw blood. Injuries that drew blood were cleaned with alcohol and treated with antibiotic

to avoid infection. Lizards that presented natural toe loss were not toe clipped.

157 Repeated annual surveys enabled us to quantify yearly frequencies of each colour

morph, while lizard recaptures during consecutive seasons allowed us to assess

159 ontogenetic variation in ventral colouration.

160 *Mating system and assortative pairing*

Podarcis muralis exhibits a polygynandrous mating system in which adult territorial 161 males patrol and aggressively defend against intruding males an area that typically 162 overlaps the home range of one or more females (Boag 1973; Barbault and Mou 1988; 163 Edsman 1990; Oppliger et al. 2007; our own unpublished observations). As a result, 164 165 during the breeding season males mate repeatedly with females within their territory, and most copulations occur between members of established pairs that associate in 166 time and space (e.g. basking together, sharing refuges). These observations are 167 consistent with reports of male mate guarding in other lacertid species (e.g. Olsson 168 1993a; Marco and Pérez-Mellado 1999; Martín and López 1999), and suggest that 169 consistent spatiotemporal male-female associations during the breeding season can be 170 used as an indicator of mating pattern. 171

To examine the existence of assortative pairing, we conducted field surveys 172 using two complementary sampling procedures. First, we used data on male-female 173 174 spatial association obtained from individually identified, marked adult animals during systematic field observations conducted during the first five years at our main study 175 site. For these individuals, spatiotemporal association and interactions (e.g. 176 copulations) among males and females were well documented. Second, in 2010 and 177 2011, we supplemented longitudinal data by sampling pairs of unmarked lizards 178 observed during independent transects in areas adjacent (> 500 m) to our main study 179

site. In this second type of surveys, we used a conservative criterion to record male-180 female pairs based on previous observations of individually-identified animals: 1) we 181 only recorded pairs in which both lizards could be unambiguously assigned to a 182 morph, 2) pairs involving subadult individuals or young adults were discarded (to 183 avoid the inclusion of lizards without a fully developed colouration; see results), and 184 3) pairs in which lizards were farther than one meter apart were also discarded. To 185 avoid pseudo-replication, transects involving unmarked individuals were conducted 186 only once during the whole sampling period (2010-2011). To test for non-random 187 pairing, we performed a Fisher's exact test on the total pooled dataset. To specifically 188 189 test for assortative pairing, we used a binomial test comparing the proportion of 190 homomorphic and heteromorphic pairs. Finally, we performed a Spearman correlation analysis with male and female SVL measurements from pairs of marked lizards to test 191 the hypothesis that the pairing system is driven by size-assortative mating. 192

193 **Results**

194 *Objective colour morph characterization*

We obtained reflectance spectra from a total of 249 adult males and 103 females with 195 pure-morph phenotypes (W, Y, O). Intermediate morphs (WO and YO) were 196 discarded for spectrophotometric measurements because the patches of their throat 197 mosaic were often too small to allow consistent measuring (see also Calsbeek et al. 198 2010). Measurement of a subsample of intermediate individuals exhibiting several 199 adjacent scales of the same colour (and thus large enough to allow consistent 200 measurement) confirmed that the colours present in intermediate morphs had the same 201 202 spectral properties as those of uniformly coloured animals (Figure S1 in sup. mat.).

Reflectance spectra from the throats of pure white, yellow and orange adult 203 204 males and females indicate that colouration does not vary continuously (Figure 1), which supports our eye-based classification into three discrete colour morphs. 205 Differences among ventral colourations are explained by variation in chroma between 206 400 and 600 nm, which gives rise to three discrete spectral steps (see error bars 207 associated with spectra in this range in Figure 1 and distribution of MC among morphs 208 209 in Figure 2). We found colour morphs to differ in colour variables, but sex and the interaction between morph and sex were not significant (Table 1 summarises Q, MC, 210 H and R₅₀ values for the three pure morphs). Finally, we found no differences among 211 212 morphs in BCI and only males differed slightly in SVL (Table 2). Using post-hoc analyses, we found O males to be larger than W males (P = 0.026), but we did not find 213 significant differences between W and Y and O and Y males (P = 0.44 and P = 0.61, 214 respectively). 215

216 Colour morph development and polymorphism stability

Data obtained from lizards recaptured during 2-3 consecutive reproductive seasons 217 (i.e. 28 females and 44 males) indicate that lizards develop their full adult colour by 218 219 the time they reach a size of 56 mm SVL, which we used as a criterion to exclude individuals without a fully-developed adult colouration. All juveniles sampled were 220 white ventrally. As they grew, some juveniles retained their white colouration as 221 adults, while others developed light yellow pigmentation or a series of isolated orange 222 scales. Subsequently, light-yellow individuals acquired an intense yellow colouration 223 and, in some cases, developed some orange scales (YO intermediate morph). In turn, 224 individuals with orange scales either remained white and orange (WO intermediate 225 morph), changed their white scales to yellow scales (YO) or, more often, developed an 226

overall orange colouration (Figure 3). No individuals examined more than once as adults underwent a colour change, suggesting that colour morphs are stable through adulthood (N = 30). At the population level, the polymorphism remained stable over time with only minor fluctuations in the relative frequency of the different morphs (Figure 4).

232 Assortative pairing

Figure 5 shows the frequency of scored pairs sorted by colour (40 independent pairs of 233 marked lizards and 77 pairs of un-marked lizards). From the subset of individually 234 identified lizards, 26 males were observed in association with females in more than 235 236 one occasion. Seventeen of these males paired with a single female. In all the remaining cases males were seen repeatedly with the same female and only 237 sporadically with other females. In these cases, we considered the female with which 238 the male paired consistently as his mate. Pairs observed repeatedly were tallied only 239 240 once.

Pairing was not random with respect to male and female colour morphs 241 (Fisher's test, P < 0.001), pure-morph homomorphic pairs of lizards being 242 significantly more common than heteromorphic pairs (Binomial test, N = 94, k = 69, P 243 < 0.001). Due to small sample sizes (N = 23), pairings involving intermediate morphs 244 were not included in the analysis (see Table S1 in sup. mat. for raw data). Although 245 SVL measurements were only available for a subset of individually identified lizards, 246 the correlation between male and female SVL within mated pairs was not significant 247 $(N = 17, \rho = 0.10, P = 0.69).$ 248

249 Discussion

To our knowledge, no study to date has examined colour-polymorphic assortative 250 251 mating in *Podarcis* or in other lizards, although a recent study by Huyghe et al. (2010b) found indirect evidence that hinted at the existence of nonrandom mating in 252 populations of *P. melisellensis* with three pure morphs. Here, we provide behavioural 253 evidence of colour-assortative pairing in a wild population of *P. muralis*, which is 254 strongly suggestive of assortative mating in this species. Size-assortative mating 255 256 cannot explain the pattern reported in this study because we did not find size differences between female morphs (and only a marginal size difference between 257 orange and white-morph males), or a significant correlation between male and female 258 259 SVL in mated pairs (Olsson 1993b; Cooper and Vitt 1997; Shine et al. 2001). Similarly, spatial data for this population (not shown) reveal that home ranges of 260 lizards from different morphs overlap as often as those of lizards from the same 261 262 morph. Thus, morphs are not in any way spatially segregated and interact with each other on a daily basis so there are ample opportunities for inter-morph associations 263 (Font et al. 2012; G. Pérez i de Lanuza, E. Font and P. Carazo, unpub. data). 264

265 The results presented here suggest that mate choice patterns may be involved 266 in the maintenance of colour polymorphisms in *P. muralis* (Roulin and Bize 2007). In polymorphic systems driven by mate choice, disassortative mating is the mechanism 267 268 that best explains the durability of rare phenotypes (Pryke and Griffith 2007). In contrast, positive assortative mating will normally promote the reduction of gene flow 269 270 among morphs and, consequently, induce incipient reproductive isolation (Pryke and 271 Griffith 2007; Hughes et al. 2010b). Theoretical arguments, mathematical models and recent empirical evidence all suggest that the processes generating and maintaining 272 colour polymorphism tend to promote speciation (e.g. Gray and McKinnon 2007; Otto 273 et al. 2008; Hugall and Stuart-West 2012). However, the existent of social and 274

ecological constraints on mate availability may curtail the divergence of colour
morphs into separate species despite strong assortative mate preferences, in which
case assortative mating will contribute to the maintenance of the polymorphism (Pryke
2009).

279 Our own results reveal a complex situation in P. muralis, with obvious morphassortative pairing but also a considerable proportion of heteromorphic pairs (38.5%) 280 that could reflect the existence of costs and/or constraints of assortative mating (Pryke 281 282 2009), or other processes such as condition-dependent variation in mate choice (Bleav and Sinervo 2007). We hence suggest that positive assortative mating could contribute 283 to the maintenance of colour polymorphisms in this species even though it is unlikely 284 that this is the only evolutionary process involved (Pryke 2009). As a matter of fact, 285 colour polymorphisms are widespread in the genus *Podarcis*, which suggest that other 286 selective processes, such as negative frequency dependent selection (Sinervo and 287 Lively 1996), are probably also involved in the maintenance of colour polymorphisms. 288

The findings reported here also contribute to our understanding of the form and 289 development of lacertid colour polymorphisms. First, our data show that there is no 290 291 sexual dimorphism in the spectral characteristics of male and female morphs; the only observable sexual differences are in the distribution of ventral colorations (i.e. throat 292 and belly in males, only throat in females). Second, spectrophotometric data confirm 293 the existence of discrete colour morphs in both males and females. Available data on 294 the visual system of lacertids (Wagner 1933; Swiezawska 1950; Svoboda 1969; 295 Dücker and Rensch 1973; G. Pérez i de Lanuza and E. Font, in preparation) and other 296 diurnal lizards (Fleishman et al. 1997, 2011; Loew et al. 2002; Bowmaker et al. 2005) 297 suggest that *Podarcis* lizards most likely perceive these colour morphs as discrete 298

phenotypes. Therefore, and given that morph differences are explained by variation 299 300 within the human visible range (i.e. 400-700 nm), an eye-based classification seems to be an adequate procedure for discriminating colour morphs in this species (for a 301 discussion of the pros and cons of eye-based classifications of ventral colourations in 302 lacertids see Vercken et al. 2007, 2008; Cote et al. 2008). Our results also show that 303 colour morphs in adult *P. muralis* appear to be ontogenetically stable. All 304 developmental changes observed in throat and belly colouration affected subadult 305 and/or young adult lizards (i.e., SVL < 56 mm), so criteria used to identify adult, 306 stable morphs should be taken into account in comparisons among morphs. For 307 308 example, Sacchi et al. (2007b, 2009) classified individuals with SVL larger than 50 309 mm as adults, which could have led to overestimating the proportion of white morph lizards in Italian populations of P. muralis (Sacchi et al. 2007a; see also Calsbeek et 310 al. 2010). 311

A host of interesting questions remain to be examined in more detail by future studies. In particular, it is often assumed that colour polymorphisms in lacertids are, as in other polymorphic lizards, genetically determined (Sacchi et al. 2007a,b; Calsbeek et al. 2010; Runemark et al. 2010), but controlled matings have not been conducted to confirm the heritability of colour morphs. Therefore, a crucial goal of future research should be to decipher the genetic basis underlying colour polymorphism in this and in other polymorphic species of lacertid lizards.

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- 495 Figure 1. Throat reflectance spectra from pure *P. muralis* adult male (a) and female
- (b) morphs. W, Y and O represent white, yellow and orange morphs respectively.
- 497 Spectra from white, yellow and orange scales from intermediate white-orange and
- 498 yellow-orange morphs are similar to those depicted here (see Figure S1 in sup. mat.).
- 499 Vertical lines: error bars (± 1 SEM).
- 500 Figure 2. Distribution of medium wavelength chroma (MC) measurements (males and
- females considered together) from each pure morph. W, Y and O represent white,
- 502 yellow and orange morphs respectively.
- 503 **Figure 3.** Developmental trajectories of colour morphs in *P. muralis* reconstructed
- 504 from recapture data. Arrows indicate direction of possible changes in colouration over
- time. Black arrows represent most frequent transitions and numbers over the arrows
- indicate the number of observed individuals for each transition in the population (f =
- 507 females; m = males). Counts in transitions from juveniles to subadults/young adults
- consider one year old animals with SVL < 56 mm (which always develop their ventral
- colouration from white juveniles). Transitions from subadults/young adults to adults
- were calculated considering only individuals captured for the first time with SVL < 56
- 511 mm, and recaptured during the following spring as adults.
- Figure 4. Changes in colour morph frequencies during a six–year period. Sample sizes
 for each sex and year are provided in brackets. *N* represents the total sample size
 across the six years.
- Figure 5. Number of male-female pairs classified by colour morph from a total of 117pairs.
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Table 1. Descriptive statistics and results of statistical analyses for colour variables:522brightness (Q), medium wavelength chroma (MC), hue (H, in degrees) and spectral523location of middle point (R_{50} , in nm) for each morph and sex. W, Y and O represent524the white, yellow and orange morphs respectively. Asterisks indicate statistically525significant results (P < 0.001 in all cases). See sample sizes in Figure 1.

		Q	MC	Н	R ₅₀
	W	38926 ± 650	0.52 ± 0.00	49.50 ± 0.65	416.54 ± 2.50
nales	Y	33500 ± 726	0.46 ± 0.00	37.83 ± 0.60	496.10 ± 3.17
Ц	0	30790 ± 1079	0.40 ± 0.01	25.56 ± 0.82	532.39 ± 5.40
7	W	41174 ± 875	0.50 ± 0.00	46.38 ± 0.01	427.07 ± 4.55
males	Y	38955 ± 1380	0.47 ± 0.00	39.06 ± 0.74	488.54 ± 5.32
fe	0	30237 ± 1546	0.40 ± 0.01	26.00 ± 1.15	536.29 ± 6.58
	morph	<i>H</i> =14.70*	<i>H</i> =39.82*	<i>H</i> =38.6*	$F_{1,346} = 329*$
	sex	H = 0.01	H = 0.01	H = 0.02	$F_{1,346} = 0.57$
interaction		H = 1.43	H = 0.66	H = 0.77	$F_{1,346} = 0.12$

Table 2. Mean values of morphometric variables SVL (i.e. snout to vent length) and
BCI (i.e. body condition index), and results of comparisons among morphs. Values
were log-transformed prior to the analyses. W, Y and O represent the white, yellow
and orange morphs respectively.

	SVL	(mm)	BCI		
	females	males	females	males	
W	61.78 ± 0.73	62.57 ± 0.51	-0.16 ± 0.10	0.21 ± 0.06	
Y	63.56 ± 0.77	63.63 ± 0.51	-0.28 ± 0.15	0.18 ± 0.08	
0	63.38 ± 0.75	64.77 ± 0.57	-0.49 ± 0.17	0.02 ± 0.11	
	$F_{2,65} = 1.71$	$F_{2,168} = 3.58$	$F_{2,65} = 1.071$	$\chi^2 = 2.58$	
	<i>P</i> = 0.19	<i>P</i> = 0.03	<i>P</i> = 0.35	<i>P</i> = 0.28	









