

Giraffe coat patterns

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1 **Seeing spots: Quantifying mother-offspring similarity and assessing fitness**

2 **consequences of coat pattern traits in a wild population of giraffes (*Giraffa***

3 ***camelopardalis*)**

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11 RH: Giraffe coat patterns

12

13 **Abstract**

14 Polymorphic phenotypes of mammalian coat coloration have been important to the study of
15 genetics and evolution, but less is known about the inheritance and fitness consequences of
16 individual variation in complex coat pattern traits such as spots and stripes. Giraffe coat
17 markings are highly complex and variable and it has been hypothesized that variation in coat
18 patterns most likely affects fitness by camouflaging neonates against visually hunting
19 predators. We quantified complex coat pattern traits of wild Masai giraffes using image
20 analysis software, determined the similarity of spot pattern traits between mother and
21 offspring, and assessed whether variation in spot pattern traits was related to fitness as
22 measured by juvenile survival. The methods we described could comprise a framework for
23 objective quantification of complex mammal coat pattern traits based on photographic coat
24 pattern data. We demonstrated that some characteristics of giraffe coat spot shape were likely
25 to be heritable, as measured by mother-offspring regression. We found significant variation in
26 juvenile survival among phenotypic groups of neonates defined by multivariate clustering
27 based on spot trait measurement variables. We also found significant variation in neonatal
28 survival associated with spot size and shape covariates. Spot trait variation also may be
29 relevant to other components of fitness, such as adult survival or fecundity. These findings
30 will inform investigations into developmental and genetic architecture of complex mammal
31 coat patterns and their adaptive value.

32 **Introduction**

33 Complex color patterns such as spots and stripes are found on many animal species and these
34 phenotypic traits are hypothesized to play adaptive roles in predator and parasite evasion,
35 thermoregulation, and communication (Cott, 1940; Caro, 2005). Many foundational studies
36 of coloration using starkly different color morphs from diverse taxa such as insects
37 (Kettlewell, 1955; Wittkopp *et al.*, 2003), mice (Morse, 1978; Russell, 1985; Bennett &

38 Lamoreux, 2003), reptiles (Rosenblum *et al.*, 2004; Calsbeek *et al.*, 2008), fish (Endler, 1983;
39 Irion *et al.*, 2016), and birds (Roulin, 2004) demonstrated Mendelian inheritance and natural
40 selection, and discovered genes that cause color morph mutations (Hoekstra, 2006; Protas &
41 Patel, 2008; San-Jose & Roulin, 2017). Individual variation in a complex color pattern trait of
42 spot size was also part of the earliest work on genetics and inheritance (Wright, 1917).
43 Measuring individual variation in complex color patterns, especially detailed measurements
44 such as animal biometrics (Kuhl & Burghardt, 2013), can provide novel insight into
45 developmental and genetic architecture (Bowen & Dawson, 1977; Klingenburg, 2010; San-
46 Jose & Roulin, 2017), and the adaptive value of the patterns (Hoekstra, 2006; Allen *et al.*,
47 2010), as well as benefitting studies of behavior (Lorenz, 1937; Whitehead, 1990), and
48 population biology (Holmberg *et al.*, 2009; Lee & Bolger, 2017), and the growing field of
49 phenomics (Houle *et al.*, 2010). Some methods to robustly quantify individuals' continuous
50 variation in complex color patterns have been developed for general use (Schneider *et al.*,
51 2012; Van Belleghem *et al.*, 2018) and specific taxa such as fishes (Endler, 1980; Holmberg
52 *et al.*, 2009), butterflies (LePoul *et al.*, 2014), penguins (Sherley *et al.*, 2010), and primates
53 (Allen *et al.*, 2015). We see a need for more tools and techniques to reliably quantify
54 individual variation in complex coat pattern traits in wild populations (Eizirik *et al.*, 2010;
55 Willisch *et al.*, 2013), and studies that use quantitative genetics and demographic methods to
56 investigate heritability and adaptive significance of those traits in wild mammal populations
57 (Kruuk *et al.*, 2008; Kaelin *et al.*, 2012).

58 The coat patterns of Masai giraffes (*Giraffa camelopardalis tippelskirchii*) are
59 complex and show a high degree of individual variation (Dagg, 1968; **Figure 1**). Masai
60 giraffes spots vary in color and shape from those that are nearly round with very smooth
61 edges (low tortuousness), to extremely elliptical with incised or lobate edges (high
62 tortuousness). Giraffe skin pigmentation is uniformly dark grey (Dimond & Montagna,

63 1976), but the spots that make up their coat markings are highly variable in traits such as
64 color, roundness, and perimeter tortuousness. This variation has been used to classify
65 subspecies (Lydekker, 1904), and to reliably identify individuals because patterns do not
66 change with age (Foster, 1966; Bolger *et al.*, 2012; Dagg, 2014). Dagg (1968) first presented
67 evidence from a small zoo population that the shape, number, area, and color of spots in
68 giraffe coat patterns may be heritable, but analysis of spot traits in wild giraffes, and
69 objective measurements of spot characteristics in general have been lacking.

70 It has been hypothesized that giraffe coat patterns evolved to camouflage neonates
71 whose primary defense against predation is concealment (Langman, 1977; Mitchell &
72 Skinner, 2003); thus the most likely fitness effects from variation in coat patterns should be
73 variation in juvenile survival. Giraffe calves spend much of their time, day and night, hiding
74 in the dappled light of trees and bushes and their ability to match this background should
75 influence detection by visually hunting predators such as lions and hyenas (Endler, 1978;
76 Merilaita *et al.*, 2017). Background matching, the adaptation of an animal's coloration to
77 mimic its average background and reduce detection by visually hunting predators, is a
78 common form of camouflage (Endler, 1978; Merilaita *et al.*, 2017). Alternative hypotheses
79 about the adaptive value of giraffe coat markings include thermoregulation (Skinner &
80 Smithers, 1990), and in this social species with good visual sensory perception (Dagg, 2014;
81 VanderWaal *et al.*, 2014), markings could also facilitate individual recognition (Tibbetts &
82 Dale, 2007) and kin recognition (Beecher, 1982; Tang-Martinez, 2001).

83 Our purpose in this study was to: 1) demonstrate the use of public domain image
84 analysis software ImageJ (Schneider *et al.*, 2012) to extract patterns from image data and
85 quantify multiple aspects of the complex coat patterns of wild Masai giraffes; 2) use
86 quantitative genetics methods (parent-offspring regression) to quantify the proportion of
87 observed phenotypic variation of a trait that is shared between mother and offspring; and 3)

88 determine whether variation in complex coat pattern traits was related to a measure of fitness
89 (survival) and thereby infer the effect of natural selection (viability selection) on giraffe coat
90 patterns (Lande & Arnold, 1983; Falconer & Mackay, 1996).

91 **Materials & Methods**

92 As a general overview, our methods were to: 1) collect field data in one area of Tanzania as
93 digital images of giraffes to be used for spot pattern and survival analyses; 2) extract patterns
94 from images; 3) quantify giraffe patterns by measuring 11 spot traits; 4) use principal
95 components analysis (PCA) to reduce the dimensionality of the spot traits; 5) use mother-
96 offspring regressions to estimate the phenotypic similarity between mother and offspring of
97 the 11 spot traits and the 1st two dimensions of the PCA; 6) use k-means clustering to assign
98 giraffe calves into 4 phenotypic groups according to their spot pattern traits; 7) use capture-
99 mark-recapture analysis to estimate survival and determine whether there are fitness
100 differences among the phenotypic groups; 8) use capture-mark-recapture analysis to
101 determine whether there are fitness effects from any particular spot traits.

102 **Field Data Collection**

103 This study used data from individually identified, wild, free-ranging Masai giraffes in a 1700
104 km² sampled area within a 4400 km² region of the Tarangire Ecosystem, northern Tanzania,
105 East Africa. Data were collected as previously described in Lee *et al.* (2016a). We collected
106 data during systematic road transect sampling for photographic capture-mark-recapture
107 (PCMR). We conducted 26 daytime surveys for giraffe PCMR data between January 2012
108 and February 2016. We sampled giraffes three times per year around 1 February, 1 June, and
109 1 October near the end of every precipitation season (short rains, long rains, and dry,
110 respectively) by driving a network of fixed-route transects on single-lane dirt tracks in the
111 study area. We surveyed according to Pollock's robust design sampling framework (Pollock,
112 1982; Kendall *et al.*, 1995), with three occasions per year. Each sampling occasion was

113 composed of two sampling events during which we surveyed all transects in the study area
114 with only a few days interval between events. Each sampling occasion was separated by a 4-
115 month interval ($4.3 \text{ years} \times 3 \text{ occasions year}^{-1} \times 2 \text{ events occasion}^{-1} = 26 \text{ survey events}$).

116 During PCMR sampling events, a sample of individuals were encountered and either
117 ‘sighted’ or ‘resighted’ by slowly approaching and photographing the animal’s right side at a
118 perpendicular angle (Canon 40D and Rebel T2i cameras with Canon Ultrasonic IS 100 – 400
119 mm lens, Canon U.S.A., Inc., One Canon Park, Melville, New York, 11747, USA). We
120 identified individual giraffes using their unique and unchanging coat patterns (Foster, 1966;
121 Dagg, 2014) with the aid of pattern-recognition software Wild-ID (Bolger *et al.*, 2012). We
122 attempted to photograph every giraffe encountered, and recorded sex and age class based on
123 physical characteristics. We assigned giraffes to one of four age classes for each observation
124 based on the species’ life history characteristics and our sampling design: neonate calf (0 – 3
125 months old), older calf (4 – 11 months old), subadult (1 – 3 years old for females, 1 – 6 years
126 old for males), or adult (> 3 years for females, > 6 years for males) using a suite of physical
127 characteristics (Strauss *et al.*, 2015), and size measured with photogrammetry (Lee *et al.*,
128 2016a). In this analysis, we used only adult females and animals first sighted as neonate
129 calves.

130 All animal work was conducted according to relevant national and international
131 guidelines. This research was carried out with permission from the Tanzania Commission for
132 Science and Technology (COSTECH) Research Permit numbers 2017-163-ER-90-172, 2016-
133 146-ER-2001-31, 2015-22-ER-90-172, 2014-53-ER-90-172, 2013-103-ER-90-172, 2012-
134 175-ER-90-172, 2011-106-NA-90-172, Tanzania National Parks (TANAPA), the Tanzania
135 Wildlife Research Institute (TAWIRI). No Institutional Animal Care and Use Committee
136 (IACUC) approval was necessary because animal subjects were observed without disturbance
137 or physical contact of any kind.

138 **Quantification of Spot Patterns**

139 We extracted patterns and analysed spot traits of each animal within the shoulder and rib area
140 by cropping all images to an analysis rectangle that fit horizontally between the anterior edge
141 of the rear leg and the chest, and vertically between the back and where the skin folded
142 beneath the posterior edge of the foreleg (**Figure 1**). For color trait analysis, we used the
143 Color Histogram procedure of ImageJ (Schneider et al., 2012) on full-color images of the
144 analysis rectangle. We extracted coat patterns using ImageJ to convert full-color images of
145 the analysis rectangle to 8-bit greyscale images, then converted to bicolor (black and white)
146 using the Enhance Contrast and Threshold commands (Schneider et al., 2012). We quantified
147 10 spot trait measurements of each animal's extracted coat pattern using the Analyze
148 Particles command in ImageJ (Schneider et al., 2012). To account for differences in image
149 resolution and animal size (including age-related growth), and to obtain approximately scale-
150 invariant standard images of each animal, we set the measurement unit of each image equal to
151 the number of pixels in the height of the analysis rectangle. Therefore all measurements are in
152 giraffe units (GU), where 1 GU = height of the analysis rectangle (**Figure 1**). We excluded
153 spots cut off by the edge of the analysis rectangle to avoid the influence of incomplete spots,
154 and we also excluded spots whose area was $<0.00001 \text{ GU}^2$ to eliminate the influence of
155 speckles.

156 We characterized each animal's coat spot pattern traits within the analysis rectangle
157 using the following 11 metrics available in ImageJ: number of spots; mean spot size (area);
158 mean spot perimeter; mean angle between the primary axis of an ellipse fit over the spot and
159 the x-axis of the image; mean circularity ($4\pi \times [Area] / [Perimeter]^2$ with a value of 1.0
160 indicating a perfect circle and smaller values indicating an increasingly elongated shape);
161 mean maximum caliper (the longest distance between any two points along the spot
162 boundary, also known as Feret diameter); mean Feret angle (the angle [0 to 180 degrees] of

163 the maximum caliper); mean aspect ratio (of the spot's fitted ellipse); mean roundness ($4 \times$
164 $[Area]\pi \times [Major\ axis]^2$ or the inverse of aspect ratio); mean solidity ($[Area] / [Convex\ area]$),
165 also called tortuousness); and mode shade ($[65536 \times r] + [256 \times g] + [b]$ using RGB values
166 from color histogram from full color photos).

167 We quantified among-individual variation in spot trait values by reporting the mean,
168 SD, and coefficient of variation (CV) of each trait. We also quantified the reliability of our
169 spot pattern trait measurement technique by computing the amount of among-measurement
170 variation for the same animal made on different photos from different dates using a set of 30
171 animals with >2 images per animal. We performed a principal components analysis (PCA;
172 Hotelling, 1933) on the covariance matrix of the 10 spot trait measurements (standardized) to
173 examine the patterns of variation and covariation among the spot measurement data and to
174 compute 2 summary dimensions explaining the 10 measurements. We performed k-means
175 clustering to divide animals into 'coat pattern phenotypes,' phenotypic groups based upon
176 their spot trait characteristics (MacQueen, 1967). The optimal number of phenotypic groups
177 was determined by the gap statistic (Tibshirani *et al.*, 2001). We performed all statistical
178 operations using R (R Core Development Team, 2017).

179 **Mother-Offspring Similarity of Spot Traits**

180 The (narrow sense) heritability of a trait (symbolized h^2) is the proportion of its total
181 phenotypic variance that is additive, or available for selection to act upon. Parent-offspring
182 (PO) regression is one of the traditional quantitative genetics tools used to test for heritable
183 additive genetic variation (Falconer & Mackay, 1996). PO regression studies cannot
184 distinguish among phenotypic similarity due to genetic heritability, maternal investment, or
185 shared environmental effects, it is however one of the few methods available when
186 information on other kin relations are lacking. Pigmentation traits in mammals are known to
187 have a strong genetic basis (Bennett and Lamoreux 2003; Hoekstra 2006), supporting the

188 interpretation of PO regression as indicating a genetic component. We expect minimal non-
189 random variation due to environmental effects because the calves were all born in the same
190 area with the same vegetation communities during a relatively short time period of average
191 climate and weather with no spatial segregation by coat pattern phenotype (**Supplementary**
192 **Material Figure S1**). The animal model was not an improvement because we do not know
193 fathers, and we had no known siblings in our dataset, therefore PO regression is the most
194 appropriate tool for our estimates of heritability, with the caveat that there are potentially
195 environmental and maternal investment effects also present.

196 We identified 31 mother-calf pairs by observing extended suckling behavior. Wild
197 female giraffes very rarely suckle a calf that is not their own (Pratt and Anderson 1979). We
198 examined all identification photographs for individuals in known mother-calf pairs, and
199 selected the best-quality photograph for each animal based on focus, clarity, perpendicularity
200 to the camera, and unobstructed view of the torso.

201 We predicted spot pattern traits of a calf would be correlated with those of its mother.
202 We estimated the mother-offspring similarity for each of the 11 spot trait measurements, and
203 the first dimension generated by the PCA. When we examined the 11 individual spot traits,
204 we used the Bonferroni adjustment ($\alpha/\text{number of tests}$) to account for multiple tests and set
205 our adjusted $\alpha = 0.0045$. We performed statistical operations in R (R Core Development
206 Team, 2017).

207 **Fitness of Spot Patterns from Juvenile Survival**

208 We assembled encounter histories for 258 calves first observed as neonates for survival
209 analysis. For each calf we selected the best-quality calf-age (age <6 mo) photograph based on
210 focus, clarity, perpendicularity to the camera, and unobstructed view of the torso, and ran the
211 photographs through the ImageJ analysis to quantify each individual's coat spot traits. We
212 analysed survival using capture-mark-recapture apparent survival models that account for

213 imperfect detectability during surveys (White & Burnham, 1999). No capture-mark-recapture
214 analyses except ‘known fate’ models can discriminate between mortality and permanent
215 emigration, therefore when we speak of survival it is technically ‘apparent survival,’ but
216 during the first seasons of life we expected very few calves to emigrate from the study area,
217 and if any did emigrate permanently this effect on apparent survival should be random
218 relative to their spot pattern characteristics.

219 We estimated age-specific seasonal (4-month seasons) survival (up to 3 years old)
220 according to coat pattern phenotype groups with calves assigned to groups by k-means
221 clustering of their overall spot traits. We compared 3 models, a null model of no group effect,
222 age + group, and age \times group, to examine whether coat pattern phenotypes affected survival
223 differently at different ages. We also estimated survival as a function of individual covariates
224 of specific spot traits including linear and quadratic relationships of all 11 spot traits and the
225 first two PCA dimensions on juvenile survival to examine whether directional, disruptive, or
226 stabilizing selection was occurring (Lande & Arnold, 1983; Falconer & Mackay, 1996). To
227 determine at what age specific spot traits had the greatest effect of survival, we examined
228 survival as a function of spot traits during 3 age periods: the first season of life, first year of
229 life, and first three years of life.

230 We used Program MARK to analyse complete capture-mark-recapture encounter
231 histories of giraffes first sighted as neonates (White & Burnham, 1999). We analysed our
232 encounter histories using Pollock’s Robust Design models to estimate age-specific survival
233 (Pollock, 1982; Kendall *et al.*, 1995), and ranked models using AICc following Burnham and
234 Anderson (2002). We used weights (W) and likelihood ratio tests as the metrics for the
235 strength of evidence supporting a given model as the best description of the data (Burnham &
236 Anderson, 2002). Due to model selection uncertainty, we present model-averaged parameter
237 values and based all inferences on these model-averaged values (Burnham & Anderson,

238 2002). We considered factors to be statistically significant if the 85% confidence interval of
239 the beta coefficient did not include zero (Arnold, 2010), this is a widely accepted convention
240 for covariates of survival analyses (Arnold, 2010).

241 Based on previous analyses for this population (Lee *et al.*, 2016a, b), we constrained
242 parameters for survival (S) and temporary emigration (γ_{\square} and γ_{\square}) to be linear functions of
243 age (symbolized 'A'), and capture and recapture (c and p) were time dependent (symbolized
244 't'), so the full model was: $\{(S(A), \gamma_{\square}(A), \gamma_{\square}(A), c(t), p(t))\}$. Giraffe calf survival does not
245 vary by sex (Lee *et al.*, 2016b), so we analysed all calves together as an additional constraint
246 on the number of parameters estimated. We tested goodness-of-fit in encounter history data
247 using U-CARE (Choquet *et al.*, 2009), and we found some evidence for lack of fit ($\chi^2_{62} = 97$,
248 $P = 0.01$), but because the computed \hat{c} adjustment was < 3 ($\hat{c} = 1.5$), we felt our models fit
249 the data adequately and we did not apply a variance inflation factor (Burnham and Anderson
250 2002; Choquet *et al.*, 2009).

251 We have deposited the primary data underlying these analyses as follows: Sampling
252 locations, original data photos, and spot trait data: Dryad DOI:
253 <https://doi.org/10.5061/dryad.6514r>.

254 **Results**

255 We were able to extract patterns and quantify 11 spot traits using ImageJ, and found
256 measurements were highly reliable with low variation in measurements from different photos
257 of the same individual (**Table 1**). From our 31 mother-calf pairs, we found two spot traits,
258 circularity and solidity (tortuousness) (**Figure 2**) had significant PO slope coefficients
259 between calves and their mothers indicating similarity (**Table 1**). Neither of the first two
260 dimensions from the PCA (see below) had significant PO regression slopes.

261 The first dimension from the PCA (from 258 calves, including the 31 calves used to
262 estimate heritability) was composed primarily of spot size-related traits (perimeter, maximum

263 caliper, area, and number) such that increasing dimension 1 meant increasing spot size.
264 Dimension 1 explained 40.5% of the variance in the data (**Figure 3**). The second dimension
265 was composed primarily of spot shape traits (aspect ratio, roundness, solidity, and circularity)
266 such that increasing dimension 2 meant increasing roundness and circularity while decreasing
267 dimension 2 meant more tortuous edges and irregular shapes. Dimension 2 explained 24.0%
268 of the variation in the data (**Figure 3**).

269 Gap statistics indicated 4 phenotypic groups was the optimal number of clusters for k-
270 means clustering, but groups 1 and 2 had a large amount of overlap in PCA variable space
271 (**Figure 4**), so we also defined 3 phenotypic groups by lumping the two overlapping groups.
272 Group 1 had medium-sized circular spots, group 2 had small-sized circular and irregular
273 spots, group 3 had medium-sized irregular spots, and group 4 had large circular and irregular
274 spots (**Figures 3 & 4**). Our survival analysis of 258 calves divided into 4 phenotypic groups
275 based on their spot traits indicated that the null model was top-ranked, but AIC_c weights
276 showed there was some evidence for survival variation among the 4 phenotypic groups
277 (**Table 2**). The 3 phenotypic group model found significant differences in survival according
278 to group (**Table 2**, beta coefficient for lumped groups 1 and 2 = -0.717, 85% CI = -1.235 to -
279 0.199). Model-averaged seasonal apparent survival estimates indicated differences in survival
280 existed among phenotypic groups during the first seasons of life, but those differences were
281 greatly reduced in ages 1 and 2 years old (**Figure 5**).

282 We found two specific spot traits significantly affected survival during the first season
283 of life (number of spots and aspect ratio; $\beta_{\text{number of spots}} = -0.031$, 85% CI = -0.052 to -
284 0.009; $\beta_{\text{aspect ratio}} = -0.466$, 85% CI = -0.827 to -0.105). Both number of spots and aspect
285 ratio were negatively correlated with survival during the first season of life (**Figure 6**). No
286 other trait during any age period significantly affected juvenile survival (all beta coefficient
287 85% CIs included zero), but model selection uncertainty was high (**Table 3**).

288 **Discussion**

289 We were able to objectively and reliably quantify coat pattern traits of wild giraffes using
290 image analysis software. We demonstrated that some giraffe coat pattern traits of spot shape
291 appeared to be heritable from mother to calf, and that coat pattern phenotypes defined by spot
292 size and shape differed in fitness as measured by neonatal survival. Individual covariates of
293 spot size and shape significantly affected survival during the first 4 months of life. These
294 results support the hypothesis that giraffe spot patterns are heritable (Dagg, 1968), and affect
295 neonatal calf survival (Langman, 1977; Mitchell & Skinner, 2003). The fact that spot patterns
296 affected survival could be related to camouflage, but could also reflect an individual quality
297 effect, maternal investment, or some other environmental effect. Our methods and results add
298 to the toolbox for objective quantification of complex mammalian coat pattern traits, and
299 should be useful for taxonomic or phenotypic classifications based on photographic coat
300 pattern data.

301 Our analyses highlighted a few aspects of giraffe spots that were most likely to be
302 heritable and which seem to have the greatest adaptive significance. Circularity and solidity,
303 both descriptors of spot shape, showed the highest mother-offspring similarity. Circularity
304 describes how close the spot is to a perfect circle, and is positively correlated with the trait of
305 roundness and negatively correlated with aspect ratio. Solidity describes how smooth and
306 entire the spot edges are versus tortuous, ruffled, lobed, or incised and is positively correlated
307 with the trait of perimeter. We did not document significant similarity of any size-related spot
308 traits (number of spots, area, perimeter, and maximum caliper), but the first dimension of the
309 PCA was largely composed of size-related traits. These characteristics could form the basis
310 for quantifying spot patterns of giraffes across Africa, and gives field workers studying any
311 animal with complex color patterns a new quantitative lexicon for describing spots. However,
312 our mode shade measurement was a crude metric, and color is greatly affected by lighting

313 conditions, so we suggest standardization of photographic methods to control for lighting if
314 color is to be analyzed in future studies.

315 We found that both size and shape of spots was relevant to fitness measured as
316 juvenile survival. We observed the highest calf survival in the phenotypic group generally
317 described as large spots that were either circular or irregular. Lowest survival was in the
318 preprint doi with small and medium-sized circular spots, and small irregular spots. Both the
319 survival by phenotype analysis and the individual covariate survival analysis found that larger
320 spots and irregularly shaped spots were correlated with increased survival. It seems likely that
321 these traits enhance the background-matching of giraffe calves in the vegetation of our study
322 area (Ruxton *et al.*, 2004; Merilaita *et al.*, 2017). However, covariation in spot patterns and
323 survival could also reflect an individual quality effect, maternal investment, or some other
324 environmental effect. The relationships among giraffe spot traits and their effects on fitness
325 are clearly complex, and require additional investigations into adaptive function and genetic
326 architecture.

327 Whether or not spot traits affect juvenile survival via anti-predation camouflage, spot
328 traits may serve other adaptive functions such as thermoregulation (Skinner and Smithers
329 1990), social communication (VanderWaal *et al.*, 2014), or indicators of individual quality
330 (Ljetoff *et al.*, 2007), and thus may demonstrate associations with other components of
331 fitness, such as survivorship in older age classes or fecundity. Individual recognition, kin
332 recognition, and inbreeding avoidance also could play a role in the evolution of spot patterns
333 in giraffes (Beecher, 1982; Tibbetts & Dale, 2007; Sherman *et al.*, 1997). Different aspects of
334 spot traits may also be nonadaptive and serve no function, or spot patterns could be affected
335 by pleiotropic selection on a gene that influences multiple traits (Lamoreux *et al.*, 2010).

336 Photogrammetry to remotely measure animal traits has utilized geometric approaches
337 that estimate trait sizes using laser range finders and known focal lengths (Lyon, 1994; Lee *et*

338 al., 2016a), photographs of the traits together with a predetermined measurement unit (Ireland
339 et al., 2006; Willis et al., 2013), or lasers to project equidistant points on animals while
340 they are photographed (Bergeron, 2007). We hope the framework we have described using
341 ImageJ software to quantify spot characteristics with trait measurements from photographs
342 will prove useful to future efforts at quantifying animal markings as in animal biometry (Kuhl
343 & Burghardt, 2013). Trait measurements and cluster analysis such as we performed here
344 could also be useful to classify subspecies, phenotypes, or other groups based on variation in
345 markings, which could advance the field of phenomics for organisms with complex skin or
346 coat patterns (Houle et al., 2010).

347 **Conclusions**

348 Masai giraffe spot patterns are particularly diverse among giraffe subspecies (Dagg, 1968),
349 and there are spot patterns in northern Tanzania that bear strong similarities to other giraffe
350 subspecies elsewhere in Africa. Two recent genetic analyses of giraffe taxonomy both placed
351 Masai giraffes as their own species (Brown et al., 2007; Fennessy et al., 2016), but the lack of
352 quantitative tools to objectively analyze coat patterns for taxonomic classification may
353 underlie some of the confusion that currently exists in giraffe systematics (Bercovitch et al.,
354 2017). We expect the application of image analysis to giraffe coat patterns will provide a
355 new, robust dataset to address taxonomic and evolutionary hypotheses.

356 Patterned coats of mammals are hypothesized to be formed by two distinct processes:
357 a spatially oriented developmental mechanism that creates a species-specific pattern of skin
358 cell differentiation and a pigmentation-oriented mechanism that uses information from the
359 pre-established spatial pattern to regulate the synthesis of melanin (Eizirik et al., 2010). The
360 giraffe skin has more extensive pigmentation and wider distribution of melanocytes than most
361 other animals (Dimond & Montagna, 1976). Coat pattern variation may reflect discrete
362 polymorphisms potentially related to life-history strategies, a continuous signal related to

363 individual quality, or a combination of both. Future work on the genetics of coat patterns will
364 hopefully shed light upon the mechanisms and consequences of coat pattern variation.

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570 **Figure Captions**

571 **Figure 1 Representative images of spot patterns of mother-calf pairs of Masai giraffes**
572 **(*Giraffa camelopardalis tippelskirchii*) from the Tarangire ecosystem, Tanzania used**
573 **in this study.** The blue rectangle shows the area analysed using ImageJ to characterize
574 spot pattern traits.

575 **Figure 2 Representative spot outlines from Masai giraffes in northern Tanzania and**
576 **their corresponding circularity and solidity values.** Ranges of spot trait values from
577 258 calves are given in parentheses.

578 **Figure 3 Contributions of 10 trait measurement variables to the first 2 dimensions of the**
579 **principal components analysis of giraffe spots.** The first dimension (Dim1) was
580 composed primarily of spot size-related traits (perimeter, maximum caliper, area, and
581 number of spots), the second dimension (Dim2) was composed primarily of spot shape
582 traits (aspect ratio, roundness, solidity, and circularity).

583 **Figure 4 Results from k-means cluster analysis of giraffe spot patterns to define**
584 **phenotypic groups.** Left is gap statistic for different numbers of groups. Right is 4
585 clusters mapped in PCA space.

586 **Figure 5 Model-averaged seasonal (4 months) apparent survival estimates for coat**
587 **pattern phenotypic groups of giraffes defined by k-means clustering of their spot**
588 **pattern traits.** There was evidence for significant differences in survival among
589 phenotypic groups during the younger ages, but those differences were greatly reduced as
590 the animals approached adulthood (age 9 – 11 seasons). Error bars are ± 1 SE.

591 **Figure 6 Survival of neonatal giraffes during their first 4-months of life was negatively**
592 **correlated with number of spots (a) and aspect ratio (b).** Number of spots and aspect
593 ratio are the inversely related to spot size and roundness (the variables used when

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594 describing coat pattern phenotypic groups), respectively. Black lines are model estimates,
595 grey lines are 95% confidence intervals.

596

597

598

599 **Table 1 Summary statistics for mother-offspring regressions of spot traits of Masai giraffes in northern Tanzania.** Mean trait values, SD
 600 (standard deviation), CV (among-individuals coefficient of variation), Reliability (mean % variation in measurements from different pictures of
 601 the same individual), PO slope coefficients, F-statistics, and P values are provided. Statistically significant heritable traits are in bold.

602

	Number	Area	Perimeter	Angle	Circularity	Maximum Caliper	Feret Angle	Aspect Ratio	Roundness	Solidity	Mode Shade	PCA 1st Dimension
Mean	18.9	0.04	0.99	87.96	0.51	0.29	88.2	1.69	0.63	0.84	6924050	
SD	7.5	0.01	0.25	15.39	0.08	0.06	14.5	0.15	0.04	0.04	3930565	
CV	0.40	0.39	0.25	0.17	0.15	0.19	0.16	0.09	0.06	0.05	0.57	
Reliability	11	11	13	4	9	8	7	5	3	2	13	
PO Slope Coefficient	0.20	0.20	0.27	0.04	0.52	0.21	-0.15	0.19	0.08	0.53	0.44	0.39
PO Coefficient SE	0.23	0.21	0.18	0.20	0.16	0.21	0.15	0.18	0.17	0.17	0.22	0.21

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F 1,29	0.76	0.87	2.27	0.04	9.97	1.01	0.91	1.11	0.19	9.73	4.16	3.45
P value	0.39	0.36	0.14	0.84	0.0037	0.32	0.35	0.30	0.66	0.0041	0.05	0.07

603

604 **Table 2 Model selection results for giraffe calf survival according to phenotypic groups**
605 **defined by spot traits.** Model weights indicated some evidence for group effects on survival.
606 Notation 'A' indicates a linear trend with age. Additive models indicate groups shared a
607 common slope coefficient, but had different intercepts; multiplicative models indicated
608 groups had different intercepts and different slopes. Model structure in all cases was {S(A ...)
609 $g''(A) g'(A) p(t) c(t)$ }. Minimum AICc = 3236.38, W = AICc weight, k = number of
610 parameters.

611

Model	Δ AICc	W	k
A + 3 groups	0	0.43	36
Null (no group effect)	0.94	0.27	34
A + 4 groups	2.06	0.15	37
A \times 4 groups	3.01	0.09	40
A \times 3 groups	3.91	0.06	38

612

613

614

615

616 **Table 3 Model selection results for giraffe calf survival as a linear or quadratic function**
617 **of spot trait covariates during the first season (4 months), first year, and first 3 years of**
618 **life.** Confidence intervals of beta coefficients for two traits excluded zero (number of spots,
619 and aspect ratio), indicating evidence for significant spot trait effects on calf survival during
620 the first season of life. Model structure in all cases was $\{S(A + Covariate) g''(A) g'(A) p(t)$
621 $c(t)\}$ with covariate structure in survival. Notation 'A' indicates a linear trend with age, 't'
622 indicates time dependence. Minimum AICc = 3239.87, W = AICc weight, k = number of
623 parameters. Models comprising the top 50% cumulative W are shown.

624

Model	Δ AICc	W	k
Number of spots, 1st season	0	0.048	33
Aspect ratio, 1st season	0.44	0.039	33
Roundness ² , 1st 3 years	0.82	0.032	34
Angle ² , 1st season	0.87	0.031	34
Roundness, 1st season	0.95	0.030	33
Solidity, 1st season	1.06	0.029	33
Area ² , 1st season	1.11	0.028	34
Circularity, 1st season	1.15	0.027	33
Angle ² , 1st 3 years	1.21	0.026	34
Null model, no covariate	1.22	0.026	32

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Maximum caliper, 1st season	1.30	0.025	33
PCA dimension 1, 1st year	1.63	0.021	33
Angle, 1st 3 years	1.75	0.020	33
Solidity ² , 1st season	1.76	0.020	34
Perimeter, 1st season	1.88	0.019	33
Feret angle ² , 1st season	1.88	0.019	34
PCA dimension 2 ² , 1st year	1.90	0.019	34
Feret angle, 1st season	1.93	0.018	33
Number of spots ² , 1st season	2.06	0.017	34

625

626

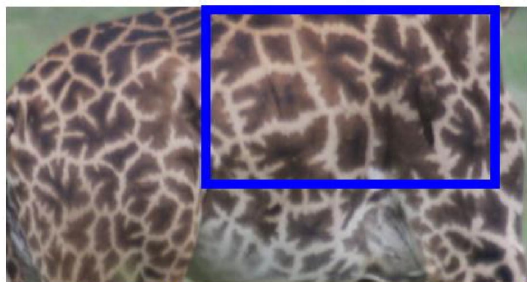
627

Pair#

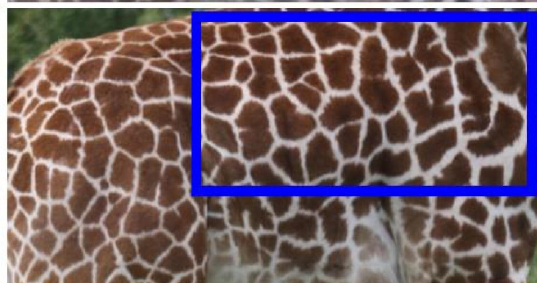
Mother

Calf

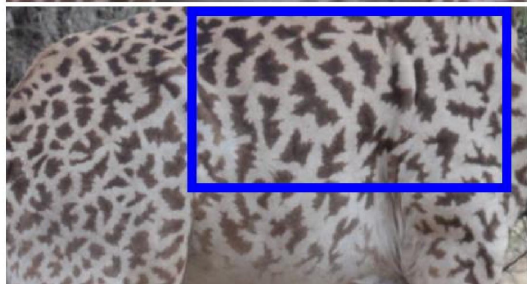
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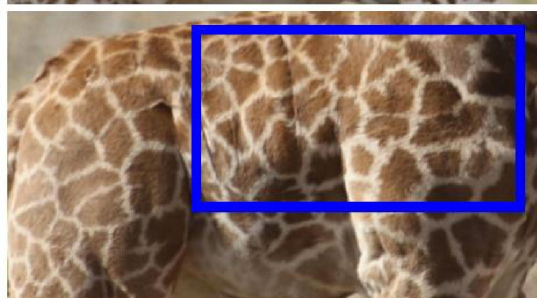
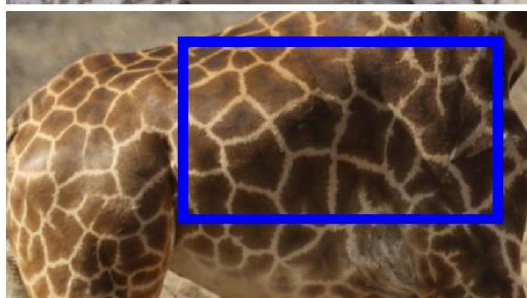
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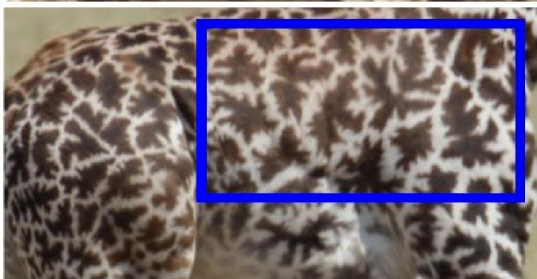
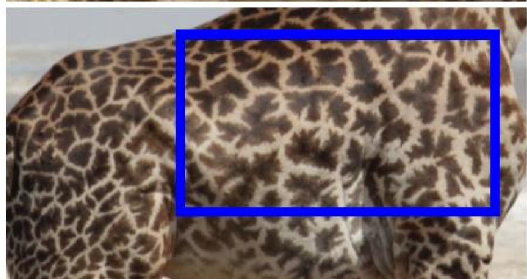
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4



5



Shape

Circularity
(0.10—0.94)

Solidity
(0.52—0.96)



0.15

0.66



0.13

0.60



0.88

0.94



0.37

0.76



0.40

0.83



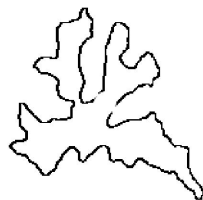
0.57

0.82



0.44

0.76



0.12

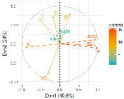
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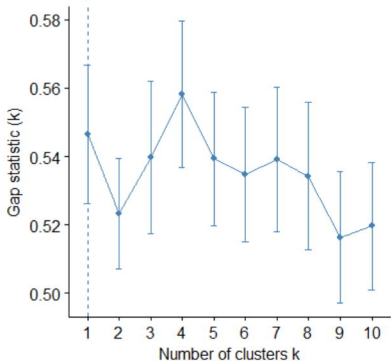
0.83

0.96

Variables - FCA



Optimal number of clusters



Partitioning Clustering Plot

