

1 **Title: *Negative impact of mild arid conditions in natural rodent populations***
2 ***revealed using markers of physiological condition in natura***

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13 **Open research statement**

14 **Data will be available from the Dryad repository (TBA)**

15 **Keywords**

16 **Aridity, blood metabolites, ecophysiology, liver, local adaptation, nutrition,**
17 ***Rhodomys dilectus dilectus*, *Rhodomys bechuanae***

18

19 **Abstract**

20 **1. Understanding how organisms respond to seasonal variations in their**
21 **environment can be a window to their potential adaptability, a classical**
22 **problem in evolutionary ecology. In the context of climate change, inducing**
23 **increased aridity and disruption of seasonality, it is crucial to study the extent**
24 **and limits of species responsiveness.**

25 **2. Here, the physiological response to food and water shortage during**
26 **seasonally dry conditions were investigated. We studied populations of two**
27 **rodent species of the genus *Rhabdomys*, one arid and one mesic, in a semi-**
28 **arid zone where their range overlap in South Africa. We measured blood**
29 **concentrations of markers of kidney and liver function, as well as body**
30 **condition, at the onset and the end of the dry season.**

31 **3. We found similar shifts in blood metabolite levels, in the semi-arid**
32 **populations of the two species, indicating malnutrition consistent with the**
33 **observed degradation of habitat quality between the start and the end of the**
34 **dry season. Furthermore, regardless of the period, differences between the two**
35 **species in blood metabolite concentrations (e.g. amylase, sodium, alkaline**
36 **phosphatase) were observed, suggesting contrasting diets and water**
37 **conservation abilities.**

38 **4. Overall, we show that, as seasonal dry conditions worsen, organisms are**
39 **increasingly affected by reduced food availability, and local adaptation to arid**
40 **conditions may provide the arid species with an advantage to cope with semi-**
41 **arid conditions. Our study suggests that even mild arid conditions could have**

42 **a negative impact and questions resilience of animals to harsher arid**
43 **conditions.**

44 **Introduction**

45 Organisms are constantly exposed to a wide range of environmental variations,
46 including short-term changes during their lifetime and longer-term changes across
47 generations (Lopez-Maury et al., 2008). In the coming decades, extreme and more
48 frequent droughts are predicted by most projections, resulting from changes in
49 precipitation patterns, increased temperature, and evaporation (Naumann et al.,
50 2018). These changes will induce great challenges to organisms. Under such
51 circumstances, divergent selection can lead to the evolution of local adaptations
52 where, in a given environment, resident phenotypes outperform nonresident ones
53 (Kawecki & Ebert, 2004).

54 As a result of climate change, organisms can be faced with warmer and drier
55 environmental conditions (Parmesan et al., 2000), likely to impact their capacity to
56 maintain homeostasis (i.e. the state of steady internal conditions allowing optimal
57 body functioning) (Davies, 2016; Fuller et al., 2016). In dry environments, food and
58 water can be scarce, generating strong selection on physiological attributes
59 maximizing energy and water availability for body function. Habitat generalists and
60 specialists' taxa may respond differently to environmental changes. Desert specialists
61 exhibit specific physiological and/or behavioral traits, resulting from genetic
62 adaptations to the environment, allowing them to continuously endure or evade
63 periods of resource restriction (Rocha et al., 2021). In contrast, generalist species
64 may seek to escape the effects of aridity through temporary avoidance (Abraham et
65 al., 2019) or plastic adjustments during unfavorable seasons (Kobbe et al., 2011).

66 Compared to mesic species, arid species have lower basal metabolic rates and
67 evaporative water loss (Muñoz-García et al., 2022), adopt a more flexible diet
68 (Tshikae et al., 2013), and cope better with prolonged droughts (Boyers et al., 2021).

69

70 Local ecological adaptation can be described in several different and complementary
71 ways, such as through its genetic basis, transcriptomic, behavioral, or physiological
72 aspects. Adaptive evolution *sensu stricto* occurs when the genetic constitution of a
73 population changes because of natural selection (Merilä & Hendry, 2013). In contrast,
74 local adaptation *sensu lato*, also considers concepts such as phenotypic plasticity
75 (the ability of a genotype to produce distinct phenotypes when exposed to different
76 environments throughout its ontogeny; Pigliucci, 2005). It is usually assessed using
77 comparative population or species fitness estimates under novel environmental
78 conditions (Webster & Reusch, 2017). The response to local environmental
79 conditions is a major factor in the generation and maintenance of biodiversity,
80 impacting population dynamics, biogeographical ranges, and species interactions
81 (Blanquart et al., 2013; Post et al., 2009). Specifically, physiological limits can drive
82 and be driven by evolution, shape species distributions and niches, and define
83 species response capacities to future climate change, directly impacting risks of
84 extinction (Somero, 2012). For example, latitudinal niches are associated with
85 adaptive variation in thermal limits in marine invertebrates (Somero, 2010).

86 Studies of different physiological systems, such as the cardiovascular (Seebacher et
87 al., 2005, Berkel & Cacan, 2022) or digestive systems (Naya et al., 2011), in
88 response to different abiotic conditions, can improve our understanding of the causal
89 mechanisms of responses of organisms to climate variation (Seebacher & Franklin,

90 2012). Biotic factors, such as sex, age, diet, strain, or breeding status also influence
91 individual responses (McClure, 1999). With a focus on influence of air temperature,
92 water availability and energy resources on an organism's physiological state,
93 research in physiological ecology has addressed how variation in physiology evolves
94 and is maintained in relation to environmental conditions (Feder & Block, 1991).

95 Analysis of blood metabolites has successfully been used to investigate kidney and
96 liver physiology of wild populations (e.g. Al-Eissa et al., 2012). The kidney and the
97 liver both contribute to homeostasis in periods of food and water scarcity; indeed, the
98 kidney plays a crucial role in maintaining osmotic balance and the liver is involved in
99 dietary metabolism. Water loss can also mechanically induce high concentrations of
100 some blood markers and impact organism's capacity to evacuate products of
101 metabolic waste, such as urea, from the blood (Ostrowski et al., 2006). Hence, dry
102 conditions can put a strain on these organs and be detrimental to survival and
103 reproduction.

104 In the Succulent Karoo of South Africa, where the average daily precipitation during
105 the dry season is 0.26 mm *versus* 0.78 mm during the wet season, the probability of
106 survival of adult *R.habdomys pumilio*, ~~another species of the same genera species of~~
107 [African striped mouse](#), were related to their physiological response at the start of the
108 dry season (Schoepf et al., 2017a). Schoepf and collaborators found higher serum
109 concentrations of albumin, glucose, potassium, and lower concentrations of globulin,
110 in animals that survived compared to those that did not survive the dry season. At the
111 peak of the dry season, individuals had lower concentrations of glucose and
112 phosphorus and higher concentrations in globulin and urea nitrogen if they survived
113 the dry season. These results strongly suggest that such markers could be good
114 indicators of individual fitness.

115 ~~The two study species, *R. bechuanae* and *R. dilectus dilectus*, have different~~
116 ~~environmental niches (du Toit et al. 2012, Meynard et al. 2012). Throughout most of~~
117 ~~its distribution, *R. d. dilectus* occurs in mesic habitats with vegetation cover and nests~~
118 ~~in dense grass, while *R. bechuanae* is found in semi-arid and arid habitats, thriving~~
119 ~~predominantly in sparsely vegetated areas and nesting in bushes (Dufour et al.,~~
120 ~~2015; Dufour et al., 2019). Moreover, differences in morphology and behavior~~
121 ~~suggest adaptation to dry conditions in *R. bechuanae* (Ganem et al., 2020; Dufour et~~
122 ~~al., 2019).~~

123 In our study, potential impact of increased aridity was addressed by investigating the
124 physiological consequences of seasonal variation in dry conditions, in two species of
125 the diurnal African striped mouse genus *Rhabdomys*. The two study species,
126 *R. bechuanae* and *R. dilectus dilectus*, have different environmental niches (du Toit et
127 al. 2012, Meynard et al. 2012). Indeed, throughout most of its distribution,
128 *R. d. dilectus* occurs in mesic habitats with dense ground vegetation cover and nests
129 in dense grass, while *R. bechuanae* is found in semi-arid and arid habitats, thriving
130 predominantly in sparsely vegetated areas and nesting in bushes (Dufour et al.,
131 2015; Dufour et al., 2019). Moreover, differences in morphology and behavior
132 suggest adaptation to dry conditions in *R. bechuanae* (Ganem et al., 2020; Dufour et
133 al., 2019). Here we address both physiological responses to seasonal dry conditions
134 and divergence ~~in-between~~ populations of the two species in the same bioclimatic
135 region. ▲

136 Indeed, at the edge of their distributions in central South Africa, the two species
137 inhabit a semi-arid region within which they occur either as parapatric or sympatric
138 populations. This semi-arid zone, the expansion and contraction of which depends
139 on land use and precipitation (Lian et al., 2021), has experienced a recent trend

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140 towards desiccation (Jury, 2021). Such drying conditions could generate additional
141 selective pressures to those already experienced by the semi-arid populations of the
142 two species. Short-term climatic patterns can also impact resource availability in this
143 region. Indeed, during La Niña years (like during this study), as part of the El Niño
144 Southern Oscillation phenomenon, there is a general association between regional
145 wetness and sea surface temperatures in the neighbouring Atlantic and Indian
146 Oceans, leading to wetter episodes throughout Southern Africa (Nicholson & Selato,
147 2000).

148

149 While many studies have harnessed physiology, genomics, and transcriptomics to
150 highlight potential adaptive interspecific or interpopulational contrasts, studies under
151 common natural environmental conditions are lacking (Rocha et al., 2021). These
152 approaches provide a mean to disentangle the roles of interpopulational or
153 interspecific variation from extrinsic environmental factors in shaping phenotypic
154 variation, taking advantage of the natural conditions experienced in the field,
155 including their complexity, with little human involvement. This study took place in a
156 natural semi-arid environment and asked how seasonal variation in dry conditions
157 influenced the physiological responses of *R.bechuanae* and *R.d.dilectus*. We
158 compared body condition, blood concentrations of markers of kidney and liver
159 functions and habitat characteristics of parapatric populations of the two species at
160 the start *versus* the end of the dry season. First, we expected that the habitat
161 available for the mice would be drier at the end compared to the onset of the dry
162 season, inducing a reduction of food and water, impairing liver and kidney function,
163 and impacting body condition. Second, assuming local adaptive plasticity, we
164 hypothesized that both species would be able to adjust their physiological responses

165 to seasonal changes in dry conditions. Third, we expected that *R. bechuanae*, having
166 evolved in arid environments, would perform better in semi-arid conditions and be
167 closer to the expected local optimal response to increased dry conditions than
168 *R. d. dilectus* (having evolved in mesic areas), as found for *Mus musculus* (Bittner et
169 al., 2021). Alternatively, if both species evolved specific adaptations to the semi-arid
170 environment, we expected no species differences (**Table 1**).

171 **Materials and Methods**

172 **Ethics statement**

173 Permits to sample and handle animals in the field were obtained from the DESTEA of
174 the Free State, the North West DEDECT and Northern Cape DENC (respectively,
175 n°202110000007994, NW 38956/04/2022, FAUNA 0389/2022). Animal handling was
176 performed under ethical clearance from the Languedoc-Roussillon ethical committee
177 for animal experimentation (n°2022041512256467 v5).

178 **Choice of sampling periods and sites**

179 We selected six study sites based on their geographical position and their aridity
180 index (AI). The AI calculation was based on Thornthwaite method (Thornthwaite,
181 1948; see **Supplementary Material** for full formula) using rainfall and temperature
182 data collected from 2010-2021 (South African Weather Service).

183
184 We selected parapatric populations of the two species in a semi-arid region of South
185 Africa ($0.2 < AI < 0.5$). In this region, a hot and wet season occurs from roughly
186 December to February, and a cool dry season from May to September. We sampled
187 six sites at the onset (May 2022) of the dry season: Barberspan Bird Sanctuary,
188 Benfontein Nature Reserve, Bloemhof Dam Nature Reserve, Gariiep Dam Nature

189 Reserve, Kalkfontein Dam Nature Reserve and Wolwespruit Nature Reserve
190 (**Figures 1 -3, Table 2**). We then resampled four of these sites at the end of the dry
191 season (September 2022). Two sites (Barberspan and Benfontein) were excluded
192 from the September sampling due to extreme weather conditions.

193 **Capture and field protocol**

194 We used small mammal PVC traps (LxHxW= 29.6x7.5x7.5 cm), baited with a mixture
195 of oats, peanut butter, salt and sunflower seeds, and added a piece of cotton wool.
196 They were placed approximately every 15 m along roughly 150 to 300 m transects.
197 Number and length of transects varied with habitat and terrain conditions. Overall,
198 trapping effort was 9688 trap nights (details in **Supplementary Tables 1&2**). Since
199 *Rhabdomys* is mostly diurnal, the traps were checked twice a day (8 AM and 3 PM).
200 On the field, all trapped striped mice were measured (body length and mass), and their
201 sex and breeding status (breeding or non-breeding) assessed based on external
202 morphological features: ~~(individuals considered as breeders presented either signs~~
203 ~~of lactation, a perforated vagina, presence of a vaginal plug, ~~or~~ scrotal testes.)~~
204 Additionally, and a 0.5 mm piece of tail was collected ~~and~~ kept in 98 % ethanol for
205 species identification. All striped mice were marked with a unique ear-tag before their
206 release at the trapping location unless they were kept for the physiological study (i.e.
207 kept in their trap with cotton wool and food).

208 Out of 903 trapped small mammals, 694 were *Rhabdomys*. For the physiological
209 study, we selected only adults, avoiding related individuals as much as possible, by
210 selecting mice trapped at least 100 m apart, except for breeding pairs (a male and a
211 female) that could be trapped in the same nest. This distance was based on data
212 obtained for *R. pumilio* indicating minimal relatedness ($R=0$ between males and

213 R=0.06 between females) at this distance (Solmsen et al. 2012). In each site,
214 trapping lasted 3 to 6 days.

215 **Habitat**

216 **Vegetation composition**

217 Earlier studies have shown inter-species differences in vegetation cover and
218 structure requirements (Dufour et al., 2015). Since habitat characteristics can vary
219 locally, we characterized the vegetation structure (i.e. grass *versus* woody
220 vegetation) at a microhabitat scale (around the trap), and at a mouse home-range
221 scale (Dufour et al., 2015). Around traps in which a *Rhabdomys* was trapped, the
222 percentage of the surface composed of dry grass, green grass, dry bushes, green
223 bushes, succulent plants, holes and uncovered surfaces was assessed within 2mx2m
224 (4m²) and the general vegetation within 10x10m (100m²) quadrats centered on the
225 trap position. For each 4m² quadrat we recorded in detail the percentage and type of
226 cover at the ground level within each of four 1x1m subunits (using a metal frame); the
227 results obtained for the four subunits were then averaged. For the 100m² quadrats
228 the assessment was made following a visual inspection at eye-level height; the
229 different cover percentages were estimated by the same observer. All together we
230 characterized 236 quadrats of each type (100m² and 4m²).

231 **Normalised Differential Vegetation Index**

232 NDVI, the satellite imagery-based index informative of ground vegetation greenness,
233 was retrieved from the Copernicus Open Access Data Hub (Copernicus Sentinel-2
234 data, 2023, calculated from 10-m resolution bands). While the two above-mentioned
235 quadrats were aimed at characterizing the structure/cover of the habitat, NDVI was

236 used as an index of habitat quality. To characterize each site and sampling session,
237 we used NDVI data available for the closest day to the beginning of a sampling
238 session, which was expected to represent the conditions experienced by the mice at
239 the time of capture.

240

241 **Blood extraction and assessment of blood metabolites concentrations**

242 In this study, a total of 273 adult mice were euthanized by means of cervical
243 dislocation on their day of capture, and a blood volume of around 100 μ L was
244 collected in lithium-heparinized tubes by cardiac puncture. The breeding status was
245 confirmed for all individuals during dissection. The whole liver, the left kidney, and the
246 skull of every individual were also collected for a complementary study. Because the
247 two species could not be distinguished visually, the spleen was also harvested from
248 each mouse and subsequently analyzed for post-hoc species identification using
249 Cytochrome Oxidase I genotyping (as described in Ganem et al. 2020).

250 Levels of albumin, alkaline phosphatase, alanine aminotransferase, amylase, total
251 bilirubin, blood urea nitrogen, total protein, globulin, glucose, calcium, phosphorus,
252 potassium, and sodium were successfully measured in the whole blood of 257
253 individuals immediately after blood collection (Abaxis VETSCAN V2 technology with
254 the Comprehensive diagnostic profile cartridge), following the manufacturer's
255 instructions. The coefficient of variation (CV%) of measured levels of these markers
256 was determined from data obtained by analyzing a given blood sample twice.

257 Samples from six adult striped mice (5 *R.bechuanae*, 1 *R.d.dilectus*) were used for
258 these calculations. For these six samples, the calculated CV equaled 5.6 ± 1.9 %,
259 which is consistent with the intra-test CV% provided in the manufacturer's instruction
260 manual for other mammal species (**Supplementary Table 3**).

261 **Data preparation and analysis**

262 All statistical tests were carried out using the R software (version 4.2.1).

263 **Habitat**

264 Vegetation composition

265 Because the distribution of the variables describing the cover and vegetation type did
266 not meet the conditions necessary for further multivariate model based analyses, we
267 performed a Principal Component Analysis (package FactoMineR version 2.9)
268 including the 7 variables measured within the 4m² quadrats and within the 100m²
269 quadrats aiming to transform ~~and, as far as possible, reduce the number of the~~
270 variables in a way that would take into account each vegetation structure parameter's
271 contribution to the overall variance. In each case, over 80 % of the variance was
272 explained by the first five Principal Components (**Supplementary Tables 4 & 5**). The
273 distribution of the coordinates of all traps on the first 5 PCs complied to the statistical
274 analysis constraints and were used as response variables of a *PERMANOVA*
275 (package *vegan*, version 2.6-4). *PERMANOVA* is a multivariate statistical inference
276 tool using permutational algorithms (Anderson, 2001). This equivalent to *MANOVA*
277 operates in a distribution-free setting and is robust to non-normality of residuals as
278 well as dispersion heterogeneity, even in slightly unbalanced designs (Anderson,
279 2013). We tested differences between sites nested within sessions (start and end of
280 dry season), with 10 000 permutations, to assess whether the sites had similar
281 enough habitat characteristics to be considered as suitable replicates in other
282 analyses.

283 NDVI

284 NDVI was averaged over a 60 m radius circular buffer centered at each trap and then
285 compared across sessions and sites using a GLM Analysis of Variance, with the site
286 nested within session. To meet statistical test prerequisites, the response variable
287 was log-transformed; the analysis was followed by Tukey multiple comparisons of
288 means post-hoc tests.

289 **Age classes**

290 Among trapped *Rhabdomys*, only adults were selected for this study based on their
291 size, general appearance (fur dullness, presence of scars), and/or breeding status.
292 Body length of trapped individuals varied between 5 and 12 cm. The body length of
293 adult individuals involved in the physiological study varied between 7.7 and 12 cm.
294 To distinguish between older and younger adults, mass was plotted against body
295 length (**Supplementary figure 6**) showing a curve defined by an allometric growth
296 equation with a steeper slope at the highest and lowest body length values. Since
297 bone growth can significantly alter the concentration of some physiological markers,
298 and growth rate varies with age/body length, we considered 4 length/age classes
299 among the adults (Class A: [7.7:9 cm[; Class B : [9:10 cm[; Class C : [10:11 cm[;
300 Class D [11 cm:]).

301 **Body condition**

302 We calculated a scaled mass index of body condition using the method described in
303 Peig and Green (2009; see **Supplementary Material** for full formula), to assess the
304 state of each individual's fat reserves (Schulte-Hostedde et al., 2005). To test
305 whether breeding status, age, habitat quality, increased dryness, and/or interspecific
306 differences influence body condition of studied individuals, linear mixed effect models

307 were computed with body condition as a response variable, site as a random factor
308 and session, species, sex, session*species*sex (including all two-way interactions),
309 breeding status (nested within sex), age class and NDVI as explanatory variables.
310 Assumptions of normality and homoscedasticity of residuals were checked using
311 diagnostic plots (*graphics* package v.3.6.2.). A Tukey multiple comparisons of means
312 post-hoc test was applied to determine which inter-level differences were driving the
313 significance of each factorial effect having more than two modalities.

314 **Physiology**

315
316 We first identified twelve outliers that we removed from further analyses. These
317 outliers were animals from which we collected smaller volume of blood samples than
318 recommended for the Vetscan or were heavily hemolyzed and flagged as such in the
319 Vetscan analysis output. In addition, as total protein values were used by the
320 Vetscan analyzer to infer the Globulin (GLOB) fraction of blood protein (with
321 $GLOB=TP-ALB$), we considered this marker as not informative and removed this
322 variable from our analyses. Finally, for 199 out of 257 samples, blood concentrations
323 of creatinine (CRE) were flagged as being below the instrument's sensitivity
324 threshold; we hence excluded CRE from subsequent analyses, as this imprecision
325 would have hindered analytical performance.

326 Due to covariance in variables such as blood physiological markers, multivariate
327 statistical techniques provide a unique insight into the main patterns and effects
328 driving variation in the data. As with habitat, multivariate normality was violated in our
329 physiological dataset, so permutational multivariate analysis of variance
330 (*PERMANOVA*) was performed. 10 000 permutations were used, with 236 blood
331 samples, 12 physiological parameters as response variables, site as a random factor,

332 and session, species, sex, breeding status (nested within sex), body condition, age
333 class, and NDVI as explanatory variables. A backwards stepwise model selection
334 was performed to retain the most parsimonious model. We then sought to identify
335 which of the 12 physiological markers drove significantly the between-group
336 differences evidenced with the *PERMANOVA*. Because in the *PERMANOVA*
337 package a dedicated function was not available, twelve post-hoc univariate linear
338 mixed model (LMM) tests were performed (followed by backwards stepwise model
339 selection), initially using the parameters of the most parsimonious *PERMANOVA*
340 model. Assumptions of normality and homoscedasticity of residuals were checked
341 using diagnostic plots. In the case of total bilirubin, a permutation test had to be
342 performed instead of the *LMM*. Through a backwards stepwise model selection
343 process. For all tests, the significance level (α) was set at 0.05.

344 **Results**

345 **Variation in habitat characteristics between species and sessions**

346 **Vegetation composition**

347 We compared the vegetation characteristics surrounding each successful trap with a
348 *PERMANOVA*, using the first 5 Principal Component coordinates (**Supplementary**
349 **Tables 4 & 5**) of each quadrat as response variable, and site and session as
350 explanatory variables. We detected significant differences in overall habitat
351 characteristics between sessions (4m² and 100m²: $p < 0.001$), and sites nested within
352 sessions (4m² and 100m²: $p < 0.001$), indicating clear site differences and seasonal
353 effects on habitat structure (**Supplementary Figures 4 & 5**). Although a
354 permutational dispersion test indicated that the latter patterns could be attributable at
355 least in part to differences in dispersion between sessions or sites (see

356 **Supplementary Material : Tables 6&7; Figures 1,2,3&4**), some general trends
357 could be inferred from the data (**Supplementary Tables 7_&_8**). Indeed, quadrats
358 from *R.d.dilectus* sites (Barberspan, Bloemhof and Wolwespruit) tended to differ
359 from feature higher proportions of grass (mean 4m²: 83.99 ± 1.12 %, mean 100m²:
360 71.42 ± 1.40 %) than *R.bechuanae* ones on Principal Components most
361 characterised by variance in grass cover, such as PC1 (mean 4 m² quadrat
362 coordinates: *R.d.dilectus* -0.68 ± 0.07 , *R.bechuanae* 0.79 ± 0.15; mean 100 m²
363 quadrat coordinates : *R.d.dilectus* -0.59 ± 0.10, *R.bechuanae* 0.64 ± 0.13) and PC2
364 (mean 4 m² coordinates : *R.d.dilectus* 0.24 ± 0.09, *R.bechuanae* -0.29 ± 0.11; mean
365 100 m² coordinates : *R.d.dilectus* -0.31 ± 0.10, *R.bechuanae* 0.34 ± 0.11) (mean
366 4m²: 54.30 ± 2.57 %, mean 100m²: 53.12 ± 1.84 %). Conversely, *R.bechuanae* sites
367 (Benfontein, Gariop and Kalkfontein) tended to feature higher proportions of bushes
368 (mean 4m²: 31.07 ± 2.37 %, mean 100m²: : 27.35 ± 1.42%) than *R.d.dilectus* sites
369 (mean 4m²: 7.33 ± 1.00 %, mean 100m²: 10.49 ± 1.03 %), or bush cover, such as
370 PC4 for 100 m² quadrats (mean coordinates : *R.d.dilectus* -0.27 ± 0.10,
371 *R.bechuanae* 0.16 ± 0.09). Furthermore, differences between sessions could be
372 observed; quadrats surveyed at the end of the dry season tended to feature more dry
373 grass (mean 4m²: 60.82 ± 2.33 %, mean 100m²: 51.99 ± 1.87 %) than at its onset
374 (mean 4m² : 50.75 ± 2.04 %, mean 100m²: 40.85 ± 1.39 %). Differences between
375 sessions could also be observed on Principal Components most characterised by
376 variance in grass cover, such as PC1 at the 100m² level (mean coordinates : onset -
377 0.11 ± 0.09, end 0.84 ± 0.07) and PC2 (mean 4m² coordinates: onset 0.27 ± 0.11,
378 end -0.38 ± 0.04; mean 100 m² coordinates: onset -0.60 ± 0.11, end 0.84 ± 0.07).

379 **NDVI**

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380 The log-transformed NDVI of sites sampled at the two sessions (2 sites per species)
381 was also compared between sites and sessions using ANOVA. Site, session and the
382 interaction between these two variables (all $p < 0.001$) contributed significantly to
383 explain the NDVI variation (**Table 3**). A post-hoc Tukey multiple comparisons of
384 means test showed that all sites had a significantly lower NDVI in September
385 compared to May (**Figure 4**), indicating a poorer state of the vegetation at the end of
386 the dry season. Furthermore, while these study locations were chosen in a region
387 with similar semi-arid bioclimatic features, Gariiep Dam and Kalkfontein (where
388 *R. bechuanae* was targeted) seemed to stand out as having a lower NDVI than all
389 other sites sampled in both May and in September, with Wolwespruit and Bloemhof
390 (where *R. d. dilectus* was targeted) having the highest NDVI (**Figure 4**). Therefore, to
391 take into account the impact of variations in NDVI between the two species trapping
392 locations we included site as a random factor in analyses addressing body and
393 physiological conditions.

394 **Variation of Body condition**

395 A linear mixed model assessed the influence of session, species, breeding status,
396 age class and NDVI on body condition, including site as a random factor. The most
397 parsimonious model included sex ($p < 0.001$), age class ($p < 0.001$), breeding status
398 nested within sex ($p < 0.01$)- and the interaction between breeding status and session
399 ($p < 0.05$); session was not significant by itself $p = 0.965$, and neither was the
400 interaction between session and sex ($p = 0.683$), while the effect of NDVI verged on
401 significance ($p = 0.067$, **Table 4**). Body condition was lower at the end of the dry
402 season than at its onset for non-breeding females compared to breeding females
403 (post-hoc pairwise t-test $p < 0.05$, **Supplementary Table 10**).

404 **Variation in physiological responses**

405 A PERMANOVA was performed using the distance matrix of 12 physiological
406 markers as response variables and session, species, sex, breeding status, body
407 condition, age class, NDVI and site (random factor) as explanatory variables
408 (**Supplementary table 11**).

409 The most parsimonious model (**Table 5**) indicated an overall difference in
410 physiological parameter concentrations between *R.bechuanae* and *R.d.dilectus*
411 ($p=0.002$), as well as an influence of body condition ($p=0.019$), breeding status
412 (nested within sex) ($p=0.028$), that differed between the two species (interaction
413 effect, $p=0.022$), and the interaction between sex and species ($p=0.020$). Session
414 ($p=0.065$), and sex ($p=0.084$) verged on overall significance and were also included
415 in the model, indicating that physiological markers overall tend to shift between the
416 start and end of the dry season, and to vary between sexes. Importantly, both
417 species showed the same pattern of variation between sessions (no significant
418 species*session interaction effect, $p=0.498$, **Supplementary Table 11**).

419 Six physiological markers showed significant differences between the start and the
420 end of the dry season (**Figure 5**). Blood concentrations of markers of nutrition and
421 liver function (albumin, potassium, alanine aminotransferase, alkaline phosphatase,
422 and total bilirubin) were lower at the end compared to the onset of the dry season
423 (**Table 4**). Glucose also showed that trend, but not significantly ($p=0.0577$). In
424 contrast, globulin levels varied in the opposite direction (higher at the end compared
425 to the onset of the dry season).

426 The levels of four markers were significantly lower in *R.bechuanae* than in
427 *R.d.dilectus* : total bilirubin, alkaline phosphatase (ALP), sodium and potassium

428 (Table 6, Figure 6). However, amylase showed the opposite pattern between the
429 species.

430 Regardless of species, the levels of glucose and ALP varied significantly with sex,
431 tending to be lower in females than in males (Supplementary figure 7).

432 Discussion

433 We asked how seasonal variation of dry conditions influenced the physiological
434 response of semi-arid parapatric populations of two closely related *Rhabdomys*
435 species that evolved under distinct environmental conditions (du Toit et al., 2012).

436 Consistent with our predictions, we found strong evidence for physiological
437 divergence between the species. We also observed a tendency for seasonal
438 variation of the physiological response in both species related to energetic resource
439 depletion at the end of the dry season, supported by the shifts in the levels of 6 blood
440 parameters between the onset and the end of the dry season.

441
442 Our study took place in a relatively wet year (La Niña). Vegetation growth and
443 senescence usually follow seasonal rainfall patterns; above-average rainfall widens
444 the temporal window for vegetation growth, improving habitat conditions at the start
445 of the typical dry season (Anyamba et al., 2002). Despite this, and consistent with our
446 predictions, milder dry conditions did alter habitat quality of our study sites (as
447 attested by shifts in NDVI and habitat structure).

448

449 Seasonal variation of physiological condition

450 Despite our study year being relatively wet, mild dry conditions impacted blood
451 marker concentrations in *R. bechuanae* and *R. d. dilectus*. These changes involved 6

452 out of 12 tested physiological markers, and included markers expected to vary in
453 periods of dietary restriction, such as albumin. Contrary to our prediction, we did not
454 find a significant interactive effect between session and species, suggesting that the
455 semi-arid populations of the two species were similarly affected by seasonal dry
456 conditions. However, a follow up study during an El Niño year may be necessary
457 before one can conclude.

458 Water and nutritional stress are expected to impact the body condition of individuals
459 (Fuller et al., 2021). As expected, body condition varied with breeding status, with
460 differences between breeding and non-breeding females heightened at the end of the
461 dry season. Additionally, several physiological markers varied significantly between
462 males and females, indicating their association with physiological processes involved
463 in reproduction. Reproduction in mammals has direct physiological costs, associated
464 with increased energy, protein and calcium demands and indirectly through
465 compensatory effects such as reduction in thermogenesis, immune function and
466 physical activity (Speakman, 2008; Rintoul & Bringham, 2014; Stawski & Rojas,
467 2016; Schoepf et al., 2017b). It follows that sex-specific differences in reproductive
468 investments are associated with contrasts in body weight, fat distribution and energy
469 metabolism (Chen et al., 2012). In support, our results indicate lower levels of blood
470 glucose, and a lower body condition in females compared with males. Moreover,
471 albumin (ALB) and alkaline phosphatase (ALP) were significantly elevated in males
472 compared to females, which could also be attributable to the pleiotropic effects of sex
473 hormones (Havill et al., 2004; Sullivan et al., 2007).

474 The two species responded similarly to increased dryness while maintaining body
475 condition, a proxy of fat reserves. Similarly, *R. pumilio* individuals tended to maintain

476 their body condition stable while reducing their energy expenditure and physical
477 activity during periods of limited food availability (Rimbach, Blanc et al., 2018;
478 Rimbach, Jäger et al., 2018). The semi-arid populations of *R.bechuanae* and
479 *R.d.dilectus* could be displaying a similar behavioral strategy.

480 Blood protein levels are generally good indicators of physiological condition (Tothova
481 et al., 2016). They comprise two distinct and major components: albumin (ALB) and
482 globulins (GLOB). In our study, ALB was significantly lower at the end than at the
483 start of the dry season. ALB is synthesized in the liver and involved in the transport of
484 bilirubin and several hormones. Blood ALB levels are less prevalent in malnourished
485 individuals (Walker et al., 1990), supporting the hypothesis of reduction of quality
486 and/or availability of food at the end of the dry season. The Nubian ibex *Capra*
487 *nubiana*, a desert mammal, also displays low ALB during the dry season (Al-Eissa et
488 al., 2012), which could reflect poor nutritional conditions, kidney dysfunction leading
489 to poor water balance and high osmolality, or liver dysfunction causing low ALB
490 synthesis. We found that GLOB was significantly higher at the end than at the start of
491 the dry season. Higher GLOB levels were also reported for *R.pumilio* individuals that
492 did not survive the dry season (Schoepf et al., 2017). GLOB is a family of molecules
493 including immunity proteins and blood protein carriers (including transferrin). An
494 increase in GLOB levels usually indicates elevation of immunoglobulins and hence
495 could suggest heightened immune activity. Greater antibody production occurs in
496 response to a higher prevalence of respiratory infections during the dry season, e.g.
497 in the Nubian goat (Abdelatif et al., 2009), and/or reduced protein nutritional status in
498 young birds (Lochmiller et al., 1993). An increase in blood GLOB concentrations may
499 also reflect high levels of transferrin (Walker et al., 1990), a major GLOB that may
500 accumulate in the blood in cases of severe iron deficiency (Kasvosve & Delanghe,

501 2002). Hence, elevated blood GLOB levels at the end of the dry season may indicate
502 either or both depletion in iron resource (transferrin) and higher vulnerability to
503 infection (immunoglobulin), both resulting from or being aggravated by reduced
504 access to food. Lower levels of alkaline phosphatase (ALP), a liver enzyme, at the
505 end of the dry season in *Rhabdomys*, is also evidence of malnutrition and deficiency
506 in essential nutrients (proteins, magnesium, zinc) (Yousef et al., 2002; Saraç &
507 Saygili, 2007; Ray et al., 2017). Alanine aminotransferase (ALT), another marker of
508 liver function, also occurred at significantly lower concentrations at the end compared
509 to the start of the dry season, indicating a reduction in liver activity (Walker et al.,
510 1990) or malnutrition (Le Couteur et al., 2010). Indeed, as environmental resources
511 become scarce, essential nutrients, such as pyridoxine (or vitamin B6), that constitute
512 some ALT coenzymes, may be less available, leading to a reduction of ALT blood
513 levels (Vespasiani-Gentilucci et al., 2018).

514
515 Blood glucose, another major physiological marker of nutrition state, was also slightly
516 diminished at the end of the dry season, albeit not significantly. Fasting or low-energy
517 feeding results in lower blood glucose levels (Jensen et al., 2013).

518
519 Acute starvation also typically causes bilirubin (TBIL) to accumulate in the blood, as
520 the enzymatic process for conversion of heme into TBIL is stimulated (Thaler et al.,
521 1972). Yet, TBIL was lower at the end compared to the onset of the dry season,
522 rejecting the starvation hypothesis. Lower TBIL concentrations at the end of the dry
523 season in *Rhabdomys* could stem from low hemoglobin counts because the body
524 produces fewer red blood cells than usual; low hemoglobin can be induced by
525 several factors, among which dietary factors such as iron deficiencies (Clark, 2008).

526 Iron deficient anemia is also supported by higher GLOB concentrations observed at
527 the end of the dry season. Infection can likewise be a cause of both decreased TBIL
528 levels (Zhao et al., 2019) and higher GLOB concentrations at the end of the dry
529 season. Finally, malnutrition and/or lack of potassium intake are among the main
530 potential causes of the reduction of its levels in the blood (Their, 1986), which could
531 explain why they were the lowest at the end of the dry season, when the available
532 resources were lowest, in our study.

533
534 Overall, our results indicate that seasonal variation of some physiological markers
535 could reflect reduced primary productivity throughout the dry season, which is
536 coherent with our observation of habitat degradation (lower NDVI) at the end of the
537 dry season. Blood concentrations of various markers seem to indicate difficulties in
538 maintaining nutritional functions and the necessity of a higher immune output, caused
539 or exacerbated by food restriction. Indeed, individuals undergoing malnutrition during
540 dry conditions are expected to show low blood glucose concentrations, and
541 deficiencies in essential nutrients, leading to pathologies such as anemia (Gordon et
542 al., 1988). Indeed, animals foraging on soils potentially poor in nutrients such as iron,
543 zinc, and magnesium are more prone to malnutrition (Graham, 1991; Gupta et al.,
544 2008). Higher GLOB, lower TBIL and lower ALP blood concentrations in our study
545 might relate to such deficiencies.

546 We did not observe seasonal variation of sodium or blood urea nitrogen, blood
547 markers of kidney function or osmoregulation. Indeed, reduced environmental
548 moisture at the end of the dry season could have caused an overall increase in blood
549 osmolality, increasing blood concentrations of most biomarkers. Still, most markers
550 exhibited a significant reduction in blood concentrations. This suggests that,

551 compared to food availability, seasonal patterns in water availability may impose a
552 milder physiological cost to the striped mice throughout the dry season in this semi-
553 arid environment during a La Niña (wet) episode.

554 **Interspecific differences**

555 *R. bechuanae* is thought to have evolved in drier environments than *R. d. dilectus*.
556 Moreover, in the semi-arid region where they co-occur, we showed that *R. bechuanae*
557 populations occupy slightly drier habitats than *R. d. dilectus*. While body condition did
558 not differ significantly between species, some interspecific contrasts support our
559 hypothesis that *R. bechuanae* may have developed a better physiological capacity to
560 cope with dry conditions than *R. d. dilectus*.

561 Among the markers exhibiting a significant difference between *R. bechuanae* and
562 *R. d. dilectus*, all but amylase showed higher blood concentrations in *R. d. dilectus*.
563 Amylase is positively correlated with digestive activity and reflects differences in
564 feeding status or diet composition between the species (Hidalgo et al., 1999). Given
565 the role of alpha-amylase in digestion, lower blood amylase levels in *R. d. dilectus*
566 than *R. bechuanae* may indicate lower digestive activity due to reduced access to
567 food, or lower starch content in the diet. Pajic et al. (2019) found a correlation
568 between dietary starch content and the number of amylase gene copies in several
569 mammalian genomes, even in different habitats and with different diets. Since
570 amylase activity and amylase gene expression are directly correlated with amylase
571 gene copy number (Arendt et al., 2014), further studies could elucidate the potential
572 genetic basis of the different levels of amylase in our two study species.

573 Sodium (NA) and potassium (K) concentrations, which depend on both dietary and
574 water intake, were higher in *R.d.dilectus* than in *R.bechuanae*. These differences,
575 coupled with higher ALP and TBIL concentrations in *R.d.dilectus* compared to
576 *R.bechuanae*, could suggest a better nutrition state for *R.d.dilectus* than
577 *R.bechuanae*. However, higher levels of amylase in *R.bechuanae* cast doubt on this
578 interpretation. Instead, lower levels of blood NA, K, ALP, and TBIL in *R.bechuanae*
579 would be more parsimoniously attributed to lower overall osmolality, perhaps due to
580 better water balance in this species. A better overall ability of *R.bechuanae* to
581 conserve body moisture could rely on physiological mechanisms such as more
582 efficient excretion or osmoregulation, or through feeding on a diet optimizing water
583 intake. We hypothesize that during the evolutionary history of *R.bechuanae*, selection
584 pressures may have favored specific strategies allowing an overall more efficient
585 water regulation compared to *R.d.dilectus*. These results, combined with the
586 reduction in the levels of K, ALP and TBIL at the end of the dry season suggest both
587 species are exposed to both water and nutrition stress.

588 We found relatively little evidence that the two species suffer differentially from lack of
589 food in their semi-arid environment during a dry season impacted by La Niña
590 phenomenon. Overall, energy levels, as indicated by blood glucose, and body
591 condition, did not vary between the species. Instead, it seems that the overall
592 differences between the two species can be accounted for by lower blood osmolality
593 in *R.bechuanae*, and on average higher blood osmolality in *R.d.dilectus*, as
594 evidenced by differences in NA and K concentrations, possibly due to overall poorer
595 water regulation. High average blood amylase levels in *R.bechuanae*, compared to
596 *R.d.dilectus* during the dry season, might indicate an interspecific difference in diet
597 composition or food intake, rather than protein-energy malnutrition. Little is known

598 about *R.bechuanae* and *R.d.dilectus* diets; the genus was characterised as a
599 generalist feeding on seeds and other plant material, and insects (Curtis & Perrin,
600 1979). Some studies showed variation in diet composition within *Rhabdomys* genus
601 associated with variation in local resources (Taylor & Green, 1976). The two species
602 could have adjusted their diet to the respective local availabilities of different food
603 resources within their home ranges. For example, during the dry season and in the
604 semi-arid zone, arthropod biomass and diversity is affected by seasonal patterns in
605 rainfall. Dalerum et al. (2017) found lower arthropod trapping success and diversity
606 during the dry season, in shrub habitats compared to any other biomes, in Benfontein
607 Game Reserve, one of our study sites. Thus, compared to seeds, the reliability of
608 arthropods as a food source during dry conditions could be lower in shrublands,
609 inhabited by *R.bechuanae* across its range and by both species in the semi-arid
610 parapathy, than in grasslands inhabited by *R.d.dilectus* in allopatry. Variation in seed
611 composition could also explain this interspecific difference in amylase concentrations.
612 Indeed, granivores prefer seeds that have higher protein content and lower
613 secondary metabolites but may be forced to take a wider range of seeds in dry areas
614 than in mesic ones (Wilmer et al., 2009). If the hypothesis of a relatively ancient
615 adaptation of *R.bechuanae* to dry habitats is true, it could be expected that it would
616 also have adjusted its diet to the specific conditions encountered in dry habitats.
617 Within this ecosystem, *R.bechuanae* could forage more efficiently compared with
618 *R.d.dilectus*, which evolved in a more mesic environment; its diet could also be richer
619 in starch (such as containing more seeds/grains than leaves or fibre). Higher starch
620 intake could help maintaining water balance, starch being the best substrate for
621 metabolic water production in a dry environment (Adolph, 1964). This is consistent

622 with the other interspecific differences revealed in this study, as diet can also
623 significantly affect osmoregulation (Sabat et al., 2009).

624

625 **Perspectives and conclusions**

626 Due to climate change, populations of many species experience changes in their
627 habitats affecting their fitness and inducing shifts in their geographical distribution
628 (Parmesan et al., 2006; Fuller et al., 2010). In some species, range shifts are
629 expected to swell edge populations, as marginal habitats grow more favourable. For
630 instance, edge populations of *Wilsonia citrina* hooded warblers have repeatedly
631 expanded their ranges due to climate change (Melles et al., 2011). In other species,
632 the quality of marginal habitats occupied by populations at the range edge may
633 decline because of reduced food availability or nutrient richness, making these
634 populations particularly vulnerable to climate change and inducing range contraction.
635 For example, the koala *Phascolarctos cinereus* experiences declines in population
636 sizes and habitat contraction due to droughts (Seabrook et al., 2011). As such,
637 depending on their phenotypic plasticity, edge populations may hold the key to the
638 survival of a species in a changing environment (Wu & Seebacher, 2022, Usui et al.,
639 2023). *R.d.dilectus* edge populations may experience the same situation, as the
640 example above, in the semi-arid zone, an hypothesis that could be tested addressing
641 these populations dynamics, compared to *R.bechuanae* semi-arid populations.

642

643 In this study, we found that two closely related species with different ecological
644 characteristics displayed similar seasonal negative shifts in blood concentrations of
645 markers of nutrition and liver function in spite of a relatively mild seasonal drying of
646 their habitats. We hence expect the physiological impact of harsher dry conditions, to

647 affect the individuals' fitness even more dramatically. The fact that the habitat of
648 *R.bechuanae* studied populations was the driest at both time points, combined with
649 our findings concerning their overall better water balance and nutrition than
650 *R.d.dilectus*, point towards *R.bechuanae* coping better in the semi-arid environment
651 during the dry season. Though adaptation *per se* cannot be inferred, this comparative
652 study considering populations of the two species in the same biogeographic zone
653 adds to the growing evidence that these cryptic species of striped mice have
654 diverged phenotypically. Overall, these findings show the ecological relevance of
655 physiological approaches in investigating the effect of climatic variation on organisms
656 and question resilience of core populations of arid species and edge populations of
657 mesic species in the face of predicted enhanced aridification.

658

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668 **Author contributions**

669 G.G. conceived the idea and designed methodology with input from H.K; G.G., H.K.,
670 P.C., collected the data; H.K. analyzed the data with input from G.G.; H.K. wrote the
671 first draft of the manuscript with input from G.G.; N.P. and N.A. contributed critically

672 to the manuscript, secured funding for the project with G.G. and helped with field
673 logistics. All authors approved the final version of the manuscript.

674 **Conflict of Interest statement**

675 The authors declare they have no conflict of interest.

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995 **Tables**

996 **Table 1: Summary of main ecological characteristics of *R.bechuanae* and**

997 ***R.d.dilectus* and of our predictions on their physiological response during the**

998 **dry season**

ECOLOGICAL CHARACTERISTICS	<i>R.BECHUANA</i> E	<i>R.D.DILECTUS</i>
ENVIRONMENTAL NICHE	Arid to semi-arid (Ganem et al., 2020)	Mesic to semi-arid (Ganem et al., 2020)
PREFERRED HABITAT	Bushy patches (Dufour et al., 2015)	Continuous cover / grassland (Dufour et al., 2015)
SOCIAL STRUCTURE	Group-living (Dufour et al., 2019)	Solitary / opportunistic group-living (Dufour et al., 2019)
SPECIES-SPECIFIC RESPONSE TO SEASONAL VARIATIONS OF DRY CONDITIONS: PREDICTIONS		
BODY CONDITION	↘↘ or ↘*	↘↘
NUTRITIONAL METABOLISM	↘↘ or ↘*	↘↘
LIVER ACTIVITY	↘↘ or ↘*	↘↘
WATER BALANCE	↘↘ or ↘*	↘↘
SENSITIVITY TO INFECTION	↗↗ or ↗*	↗↗
SPECIES DIFFERENCES: PREDICTIONS		
BODY CONDITION	+	-
LIVER ACTIVITY	+	-
WATER BALANCE	+	-
SENSITIVITY TO INFECTION	-	+

999

1000 *Adaptation to harsher arid conditions could have enabled *R.bechuanae* to withstand growing aridity

1001 without suffering as many adverse effects as *R.d.dilectus* in the semi-arid zone. Alternatively, semi-

1002 arid populations of both species could both have developed local adaptation to dry conditions.

1003

1004 ***Table 2: Geographical position and Aridity Index of sampled sites for each***
 1005 ***species***

SPECIES	SAMPLED SITES	GEOGRAPHICAL COORDINATES	ARIDITY INDEX*
<i>R.bechuanae</i>	Gariiep Dam Nature Reserve	E25.55537; S30.57589	0.42 ± 0.07
	Kalkfontein Dam Nature Reserve	E25.28031; S29.52058	0.54 ± 0.07
	Benfontein Nature Reserve**	E24.81824; S28.82747	0.39 ± 0.07
<i>R.d.dilectus</i>	Bloemhof Dam Nature Reserve	E25.65144; S27.66063	0.35 ± 0.06
	Barberspan Bird Sanctuary**	E25.58354; S26.56538	0.39 ± 0.05
	Wolwespruit Nature Reserve	E26.26326; S27.41418	0.53 ± 0.10

1006 *Average Aridity index (see **Supplementary material** for formula) ± Standard Error of
 1007 the Mean, based on climate data collected from 2010 to 2021 (South African Weather
 1008 Service)

1009 ** Sampled only in May 2022 (dry season onset).

1010

1011 **Table 3: Results of the Analysis of Variance (ANOVA) test addressing variation**
1012 **in NDVI**

Variable	Df	R ²	F	P
Session	1	0.445	442.757	< 2.2*10⁻¹⁶
Site	3	0.348	115.385	< 2.2*10⁻¹⁶
Session*Site	3	0.021	7.113	1.512*10⁻⁴
Residual	229	0.185		

1013

1014

1015 ***Table 4: Results of the Analysis of Variance (ANOVA) test addressing variation***
1016 ***in body condition***

Variable	Df	R ²	F	P
Session	1	1.271*10 ⁻⁵	0.0019	0.965
Sex	1	0.085	25.163	1.072*10⁻⁶
NDVI	1	8.007*10 ⁻⁴	0.239	0.626
Age class	3	0.091	8.990	1.201*10⁻⁵
Breeding status	2	0.041	6.067	0.003
Session*Sex	1	5.592*10 ⁻⁴	0.167	0.683
Session*Breeding status	2	0.0299	4.444	0.012
Residual	229	0.753		

1017

1018

1019 **Table 5: Summary of the results of the best fitting Permutational multivariate**
 1020 **analysis of variance (PERMANOVA) model, addressing variation in**
 1021 **physiological response (site= random factor)**

1022

Variable	Df	R ²	Pseudo-F	P
Session	1	0.011	2.882	0.067
Species	1	0.042	11.070	0.003
Sex	1	0.010	2.748	0.087
Breeding status	2	0.033	4.320	0.011
Species * Sex	1	0.016	4.232	0.029
Species * Breeding status	2	0.026	3.413	0.037
Residual	227	0.862		

1023

1024

1025 **Table 6: Summary of the results of univariate mixed models addressing**
 1026 **variation in responses of each of the 12 studied physiological variables (best**
 1027 **fitting models).**

Physiological marker	Variable(s) kept in Best Model and their significance	adjusted R ²	F	df Model	df Residual	P (model)
Albumin	Session^{***}, Sex[*]	0.098	13.72	2	233	2.338*10 ⁻⁶
Alkaline phosphatase	Session^{***}, Species⁺, Sex^{***}, Breeding status⁺	0.260	17.43	5	229	< 1.281*10 ⁻¹⁴
Alanine aminotransferase	Session⁺, Species, Sex, Breeding status, Species*Sex, Species*Breeding Status	0.060	2.884	8	227	0.004
Amylase	Species^{***}, Sex, Breeding status, Species*Sex, Species*Breeding Status	0.185	8.603	7	228	3.34*10 ⁻⁹
Calcium	Species	0.009	3.095	1	234	0.080
Globulin	Session^{***}	0.291	97.44	1	234	< 2.2*10 ⁻¹⁶
Glucose	Session ⁺ , Sex^{**}	0.079	11.14	2	233	2.397 *10 ⁻⁵
Potassium	Session⁺, Species⁺	0.068	9.569	2	233	1.014*10 ⁻⁴
Sodium	Species^{***}, Sex	0.092	12.87	2	233	4.983*10 ⁻⁶
Blood Urea Nitrogen	<i>Null Random effect model</i>	0.072	/	/	/	/
Phosphorus	Species, Sex, Breeding status⁺ + random	0.248	/	/	/	(χ ² test vs. null model) 3.471*10 ⁻⁷

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 1030 *Note: The full models, before backwards elimination, included session, species, body condition, sex,*
 1031 *breeding status, sex*species and breeding status*species as variables, with breeding status nested*
 1032 *within sex and site as a partially-crossed random factor nested within species.*

1033 **Bold:** significant effects ($p < 0.05$) following Bonferroni correction with:

1034 *****:** $p < 0.001$; ****:** $0.001 < p < 0.01$; ***:** $0.01 < p < 0.05$; **+**: $0.05 < p < 0.1$

1035 **Figure captions**

1036 **Figure 1: Map of known occurrences of 6 main clades of *Rhabdomys***
1037 **(translucent dots) and locations sampled in this study (full colour dots) in**
1038 **southern Africa.** Based on published data and unpublished data (origin details are
1039 available in doi: 0d1b3414-7e2a-11ea-a38d-00163e26bfb0). Base map: World
1040 Topographic Map Esri Standard, Aridity Index layer was computed from a 0.5° global
1041 grid, using data from the Version 3 of the Global Aridity Index and Potential
1042 Evapotranspiration Database (Zomer et al., 2022).

1043 **Figure 2: Graphical summary of organization and composition of data used for**
1044 **analysis in this study.** Figures between brackets indicate number of modalities or
1045 types of categories included in the factor.

1046 **Figure 3: Total monthly precipitation for all sampled sites as recorded in 2022.**

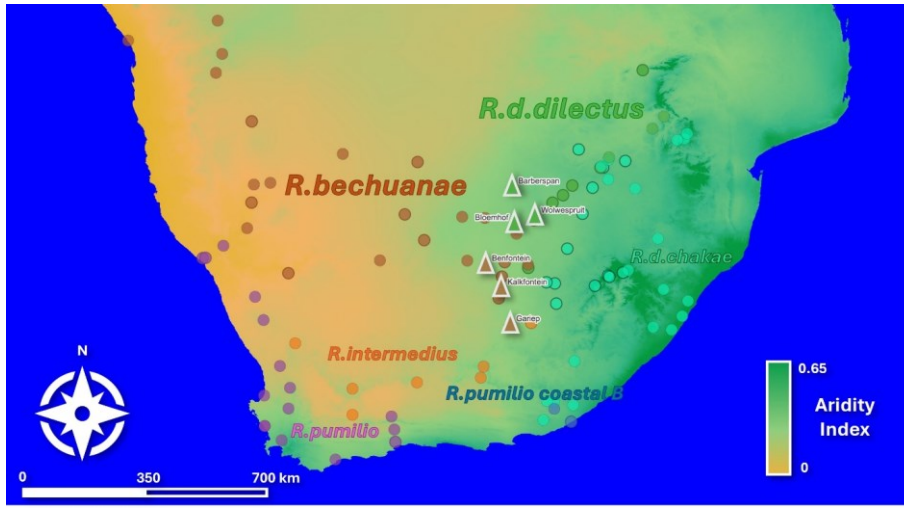
1047 **Figure 4: Average Normalized Differential Vegetation Index calculated within a**
1048 **buffer circle of 60m radius around each successful trap per site and session.**
1049 NDVI was retrieved from the Copernicus Open Access Data Hub (Copernicus
1050 Sentinel-2 data [2023]). For each site and session, we retained NDVI data available
1051 for the closest day to the beginning of a sampling session, expected to represent the
1052 conditions experienced by the mice at the time of capture.

1053 **Figure 5: Significant shifts in blood physiological marker concentrations**
1054 **between the onset and the end of the dry season (mean ± SE)**

1055 **Figure 6: Significant interspecific differences involving four blood**
1056 **physiological marker concentrations (mean ± SE)**

1057

1058 **Figures**



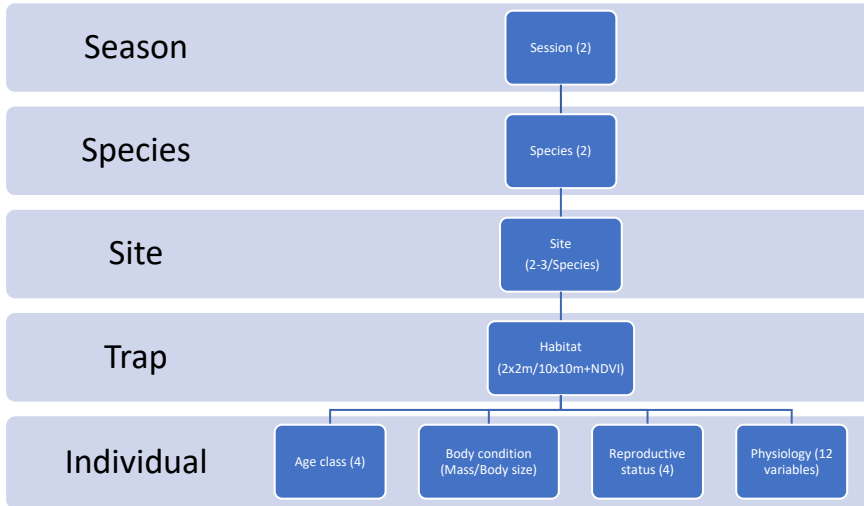
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Figure 1: Map of known occurrences of the 6 main clades of Rhabdomys (dots) and locations sampled in this study (triangles) in southern Africa

1061 *Based on published data and unpublished data (origin details are available in doi:*
1062 *0d1b3414-7e2a-11ea-a38d-00163e26bfb0). Base map: World Topographic Map Esri*
1063 *Standard, Aridity Index layer was computed from a 0.5° global grid, using data from*
1064 *Version 3 of the Global Aridity Index and Potential Evapotranspiration Database*
1065 *(Zomer et al.,2022).*

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Figure 2: Graphical summary of organisation and composition of data used for

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analysis in this study. Figures between brackets indicate number of modalities or types

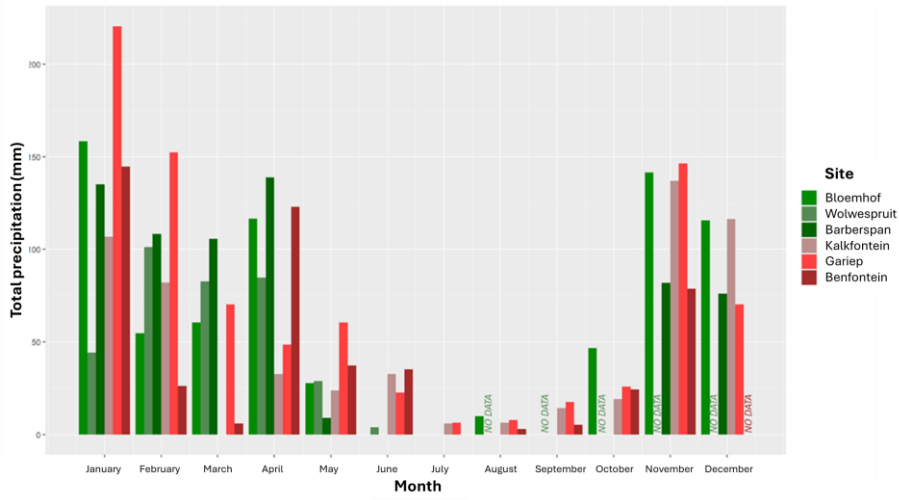
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of categories included in the factor.

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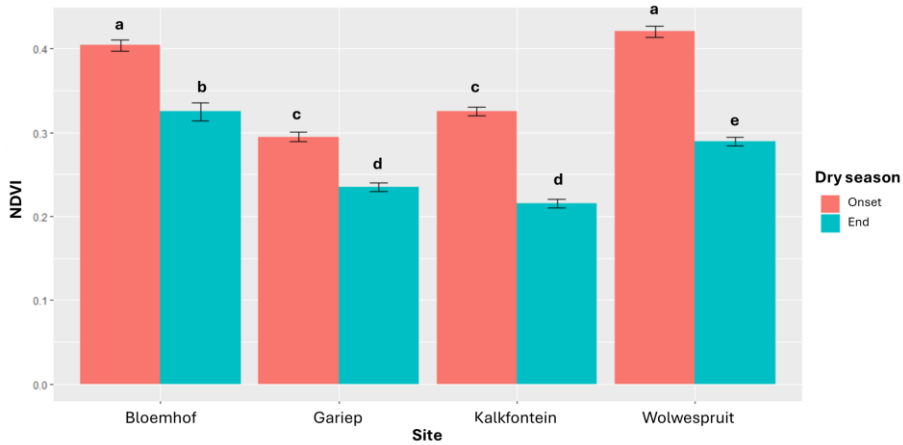
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1075 **Figure 3: Total monthly precipitation for all sampled sites as recorded in 2022.**

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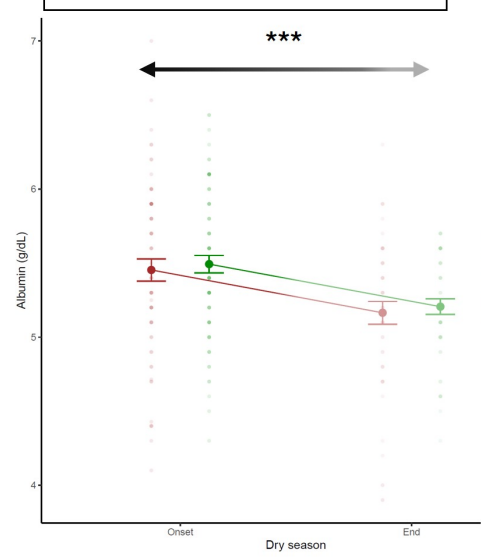
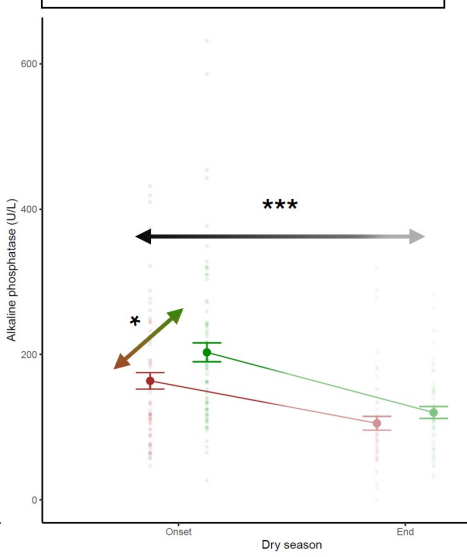
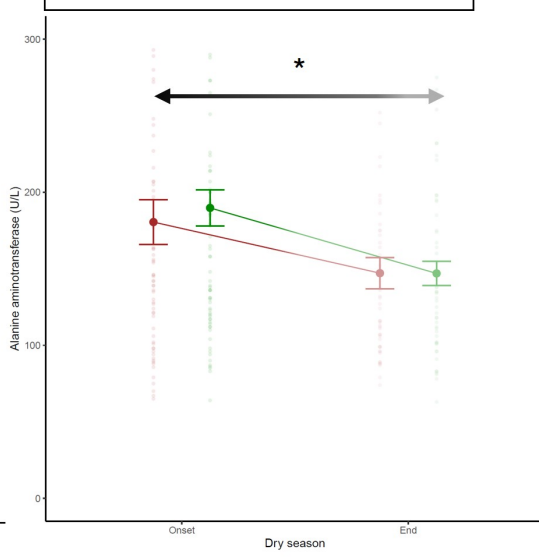
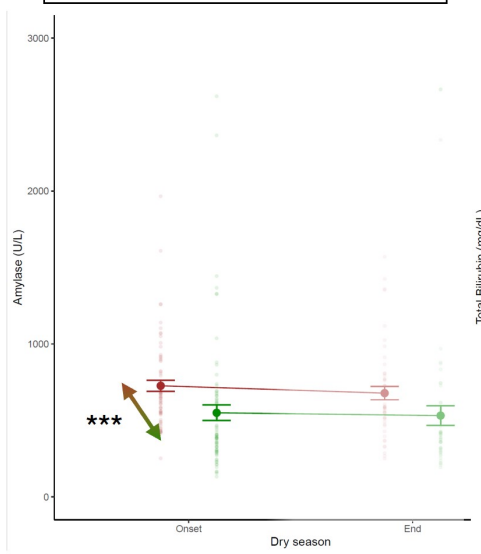
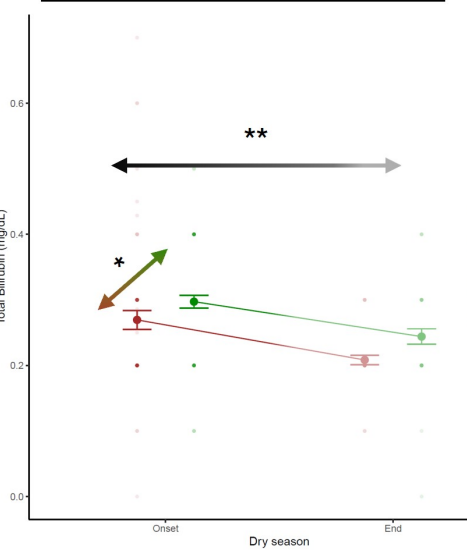
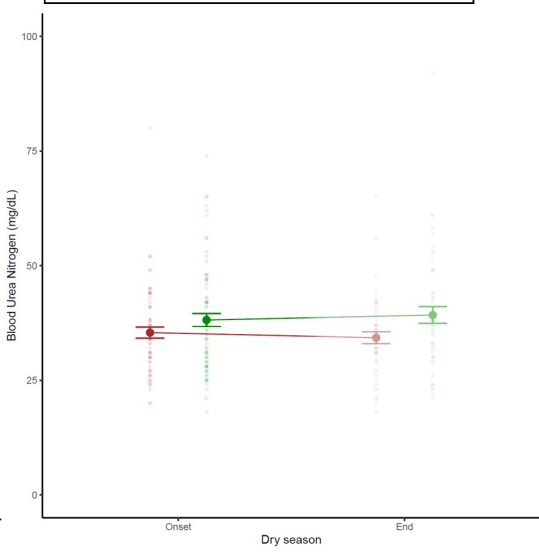
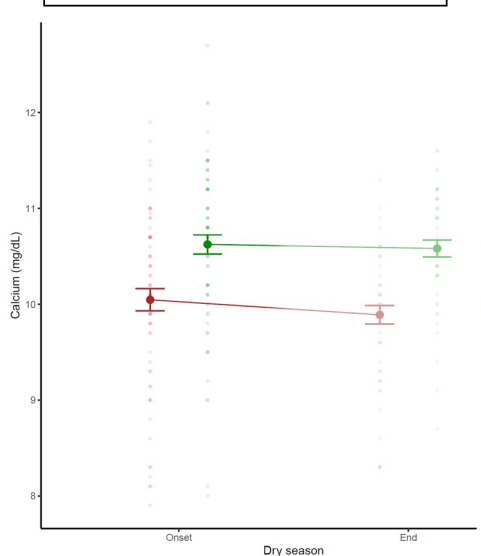
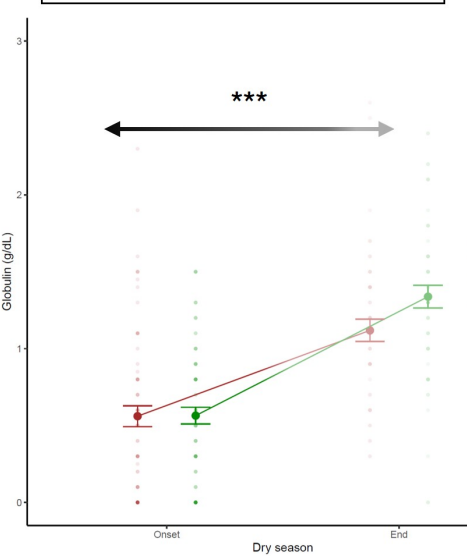
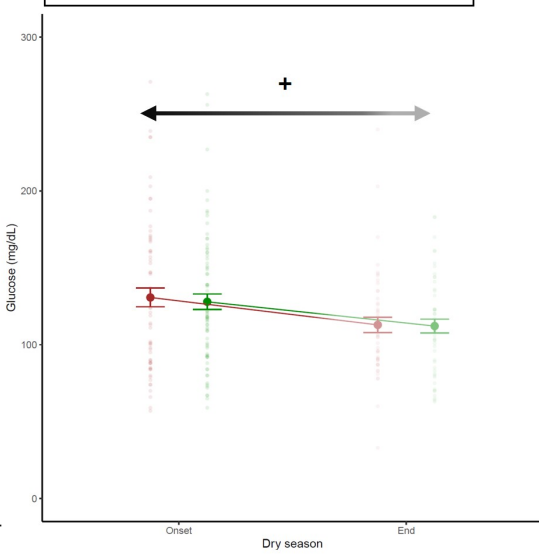
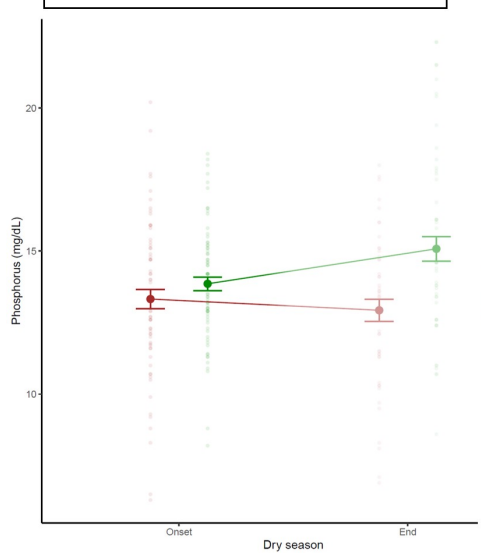
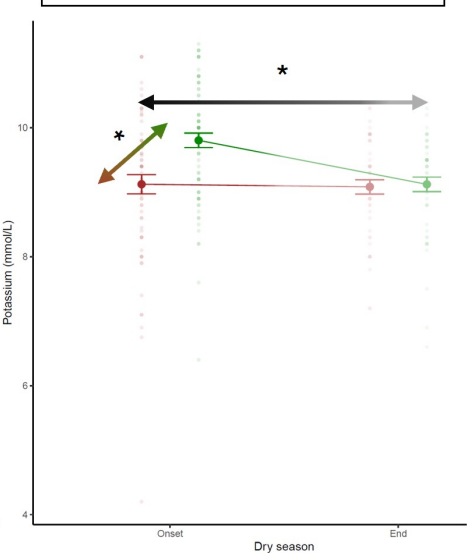
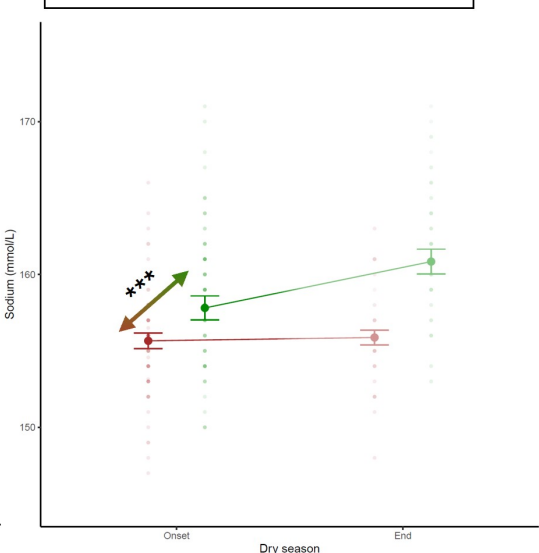
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1078 ***Figure 4: Average Normalized Differential Vegetation Index calculated within a***
 1079 ***buffer circle of 60m radius around each successful trap per site and session.***

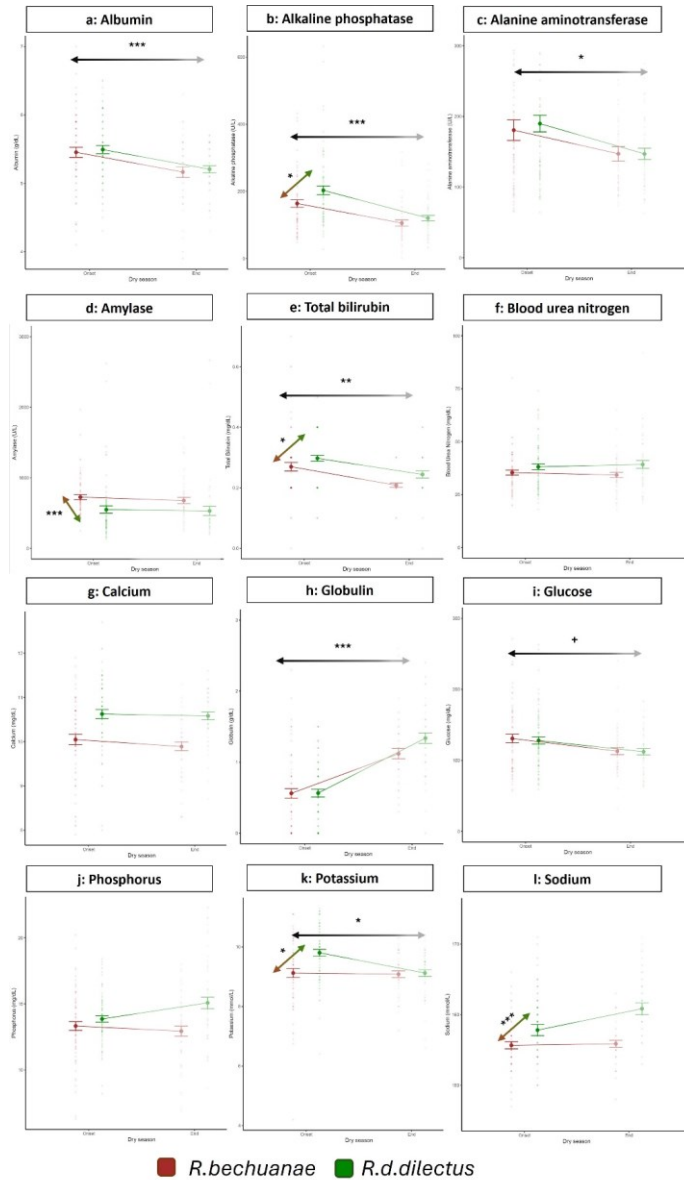
1080 *Different letters above the bars indicate statistically significant difference at $p < 0.05$.*

1081 *NDVI was retrieved from the Copernicus Open Access Data Hub (Copernicus*
 1082 *Sentinel-2 data [2023]). For each site and session, we retained NDVI data available*
 1083 *for the closest day to the beginning of a sampling session, expected to represent the*
 1084 *conditions experienced by the mice at the time of capture.*

1085

a: Albumin**b: Alkaline phosphatase****c: Alanine aminotransferase****d: Amylase****e: Total bilirubin****f: Blood urea nitrogen****g: Calcium****h: Globulin****i: Glucose****j: Phosphorus****k: Potassium****l: Sodium**

■ *R. bechuanae* ■ *R. d. dilectus*



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Figure 5 : Levels of 12 physiological markers at the onset and end of the dry season in *R.bechuanae* and *R.d.dilectus*. Arrows indicate significant effects ($p < 0.05$) between the onset and the end of the dry season (black to grey gradient) or between the two species (brown to green gradient) following Bonferroni correction, with;

*****: $p < 0.001$; **: $0.001 < p < 0.01$; *: $0.01 < p < 0.05$; +: $0.05 < p < 0.1$**

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