

1 **New insights into morphological adaptation in common mole-rats**
2 **(*Cryptomys hottentotus hottentotus*) along an aridity gradient**

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16
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26 **Abstract**

27 Morphological adaptation is the change in the form of an organism that benefits the individual
28 in its current habitat. Mole-rats (family Bathyergidae), despite being subterranean, are
29 impacted by both local and broad-scale environmental conditions that occur above ground.
30 Common mole-rats (*Cryptomys hottentotus hottentotus*) present an ideal mammalian model
31 system for the study of morphological variation in response to ecology, as this species is found
32 along an aridity gradient and thus can be sampled from geographically non-overlapping
33 populations of the same species along an environmental longitudinal cline. Using the mass of
34 five internal organs, ten skeletal measurements and 3D morphometric analyses of skulls, we
35 assessed the morphology of wild non-breeding individuals from five common mole-rat
36 populations in South Africa. We found that the body mass and mean relative mass of the
37 spleen and kidneys in arid populations was larger, and individuals from arid regions possessed
38 shorter legs and larger inter-shoulder widths compared to individuals from mesic regions.
39 Additionally, arid populations demonstrated greater skull depth, and shape change of
40 features such as angular processes of the lower jaw than mesic individuals, indicating that
41 these distinct geographic populations show differences corresponding to the aridity gradient,
42 potentially in response to environmental factors such as the variation in food sources found
43 between different habitats, in addition to different soil compositions found in the different
44 regions. Arid populations potentially require a stronger jaw and neck musculature associated
45 with mastication to chew xeric-adapted plants and to dig through hard soil types, whereas
46 mesic populations excavate through soft, looser soil and may make use of their front limbs to
47 aid the movement of soils when digging. Aridity influences the morphology of this species and
48 could indicate the impact of environmental changes on speciation and mammalian skull
49 morphology.

50 **Introduction**

51 Morphological adaptation is the change in the form and structural features of an organism
52 that are beneficial in its current habitat (Millien et al., 2006). These morphological adaptations
53 can be in relation to the physical environment, and can aid ecological, biological, physiological
54 and behavioural processes. Understanding morphological changes is integral in identifying
55 how both individuals and species respond to changes in external pressures, both in terms of

56 evolutionary history and in more rapid response to present changes in their environment
57 (Navas et al., 2004).

58 Morphology is influenced by external biotic (e.g., presence of predators and intra-specific
59 competition) and abiotic (e.g., water availability and temperature) adaptive forces. For
60 example, in response to predation, specific fur colours and forms may provide camouflage.
61 Populations of North American hog-nosed skunks (*Conepatus leuconotus*) show intraspecific
62 variation in colour patterns, with populations found in arid open environments having more
63 white colourations along the dorsum compared to those populations in the forested regions.
64 The pelage difference is believed to be a consequence of camouflage by the animals to
65 different habitat types, as the reduced whiteness and increased black colouration in forest
66 populations is thought to aid in camouflage in the dark understory (Ferguson et al., 2022).
67 Morphological adaptations are more pronounced in extreme environments as a result of a
68 greater drive for specific adaptive features. Several extreme hostile environments are found
69 on Earth and these often require distinct adaptive features for species to survive within these
70 ecosystems (Harris et al., 1998). An example of this is desert and xeric regions that form the
71 largest terrestrial biome, which in the early 21st Century covered 19% of land surface area
72 (Lockwood et al., 2012). These arid regions can be extremely harsh and are characterised by
73 ambient temperature extremes and seasonal or year-round paucity of water. This presents a
74 series of challenges for the organisms that live there, including the scarcity of resources,
75 desiccation, heat exposure, and frequent fluctuations in temperature (Harris et al., 1998).
76 Morphological adaptations to aridity can involve the development of exaggerated features
77 that aid in heat dissipation such as in fennec foxes (*Vulpes zerda*), found throughout the
78 Sahara, which possess large ears to aid in temperature control and heat dissipation (Geffen
79 and Girard, 2003). Coat colour and thickness can also facilitate heat loss in arid environments
80 (Stuart-Fox et al., 2017). Lighter fur morphs of springbok (*Antidorcas marsupialis*), found in
81 the arid Karoo region of South Africa, were found to have increased heat loss when compared
82 to dark fur morphs (Hetem et al., 2009). Furthermore, springboks were found to have thinner
83 fur than other species of similar sizes, which is also thought to increase heat loss.

84 A single species may exhibit a spatial distribution across different climates, and the influence
85 of the environment on morphology at the population level is a pivotal aspect of understanding
86 speciation and local adaptation (Ryding et al., 2021). There is widespread evidence of

87 population-level changes in appendage size in response to climate change, for example,
88 incorporating Allen's rule (Allen, 1877), which states that appendage size increases with
89 increasing ambient environmental temperature. In addition, intra-specific variation in body
90 size is found to differ with climate patterns. Bergmann's rule states that populations in colder
91 climates will have larger body sizes than those from warmer climes (Bergmann, 1847). Both
92 Allen's rule and Bergmann's rule have been observed in many taxa (but not all) including
93 insects, amphibians, terrestrial mammals and birds (Nudds and Oswald, 2007; Symonds and
94 Tattersall, 2010; Alho et al., 2011; Osorio-Canadas et al., 2016). n

95 Currently, morphological changes at the species level in response to aridity have yet to be
96 extensively studied, in particular, morphological changes of subterranean mammals (Ryding
97 et al., 2021). Furthermore, different environmental conditions interact, such that aspects of
98 morphology may often lead to adaptive trade-offs between evolutionary pressures and co-
99 evolved features in response to multiple environmental stresses, for example how
100 subterranean species cope with living underground, as well as living in arid desert conditions.
101 African mole-rats (Bathyergidae) are a group of subterranean rodents found across sub-
102 Saharan Africa including in a range of climatic regions from hyper-mesic to hyper-arid
103 (Bennett, and Faulkes, 2000). In addition to living in harsh arid habitats, mole-rats occupy a
104 subterranean niche, requiring adaptations to the small spaces of enclosed burrow systems
105 and associated low oxygen (Bennett and Faulkes, 2000). The ability of African mole-rats to
106 occupy these extreme niches has been explored in species such as naked mole-rats
107 (*Heterocephalus glaber*), distributed in the Horn of Africa, and Damaraland mole-rats
108 (*Fukomys damarensis*) across the Kalahari desert (Bennett and Faulkes, 2000). It has been
109 suggested that the ability of some African mole-rat species to persist and thrive in arid regions
110 of Africa is due to the adaptive benefits of group living which increases the energy allocations
111 dedicated to foraging and locating stochastically distributed food sources. This theory, laid
112 out by Jarvis et al. (1994) as the aridity food distribution hypothesis (AFDH) was further
113 developed by Spinks et al. (2000), and larger individual body mass was identified in arid
114 populations of *C. h. hottentotus* in comparison to mesic populations.

115 Both species exhibit eusocial behaviour, which is understood to be linked to aridity through
116 the aridity food distribution hypothesis (AFDH) (Jarvis et al. 1994). The AFDH posits that larger
117 numbers of individuals per colony are found in arid regions to increase the chances of finding

118 large and stochastically distributed food resources to secure sufficient energy allocations of
119 the colony dedicated to foraging (Jarvis et al., 1994).

120 African mole-rats are morphologically adapted to their fossorial lifestyle. Members of the
121 group possess cylindrical bodies and short limbs, with large front claws, that in some species
122 aid with movement through burrows (Jarvis, 1984; Stein, 2000; Bennett and Faulkes, 2000;
123 Gomes Rodrigues et al., 2023). Mole-rats are characterised by their large and powerful
124 extrabuccal incisors which they use to dig through the soil (Jarvis, 1984). As with all rodents,
125 these incisors continually grow throughout their life, being worn down from digging, thus
126 enabling repeated periods of burrowing when forming their underground tunnels (Single and
127 Dickman, 2018). Uniquely, their lips can close behind the incisors, creating a seal to prevent
128 soil from entering the mouth as they dig (Jarvis, 1984). Because of their fossorial life, all mole-
129 rat species have relatively small eyes and very poor vision, only able to detect light and dark
130 (Bennett and Faulkes, 2000; Burda, 2006). The digging strategy of *Cryptomys* species has been
131 described as chisel-tooth digging, whereby soil displacement is undertaken primarily by the
132 lower incisors, while the upper incisors anchor skull to the soil (Jarvis and Sale, 1971). Chisel-
133 digging is found in all mole-rat species (with the exception of those in the genus *Bathyergus*)
134 and is linked to skull morphological features such as increased depth of skull and large upper
135 incisor procumbence; the angle of the protrusion of the incisor from the rostrum (Lessa,
136 1990). This feature allows for a more favourable angle of the head for a stronger anchor in
137 the soil and provides a greater bite force and gape to dig through harder soil types (McIntosh
138 and Cox, 2016a; Kraus et al., 2022). Cranial morphology and digging type is thus suggested to
139 correlate with soil hardness, and thus with habitat type (Barčiová et al., 2009).

140 Common mole-rats (*Cryptomys hottentotus hottentotus*) are an ideal model species to
141 investigate morphological variation along an environmental cline, as this mole-rat subspecies
142 is found in South Africa and occupies a distribution along an aridity gradient (Spinks, 1998).
143 Unlike naked mole-rats (*Heterocephalus glaber*) common mole-rats possess fur. The retention
144 of pelage and the effects on thermoregulation are not currently understood. Common mole-
145 rats, like all African mole-rats, do not drink free-standing water but rely on their food source
146 of underground geophytes to obtain all their water requirements (Jacobs et al., 2022). Aridity
147 has been established to influence metabolism and water uptake in mammals, with species in
148 arid environments exhibiting lower metabolic rates and water loss than mesic-adapted

149 species (Tieleman et al., 2003). Oxidative stress in mesic common mole-rat populations has
150 been detected in the kidneys when compared to arid populations and the low levels of
151 oxidative stress in arid populations may infer these individuals have mechanisms to combat
152 hyperthermia and dehydration, and potentially exercise-induced damage due to the more
153 compact soil (Jacobs et al., 2022). Thus, precipitation and temperature, as well as the different
154 resulting soil structure may exert differing selection pressures on individuals found in arid
155 environments, compared to non-arid. Common mole-rat populations can, therefore, be used
156 for comparisons between arid and mesic conditions operating on the same subspecies, thus,
157 enabling the exploration of intra-specific morphological adaptations (Bennett and Faulkes,
158 2000).

159 We aimed to explore morphological elements including body mass, body size, various
160 measures of the limbs and feet as well as internal organ masses, and fur colour and thickness.
161 We predicted that arid-adapted populations would have a greater mass of organs whose
162 function relates to water storage and filtration, thus larger organ masses of kidneys and liver.
163 We predicted lighter fur colouration and reduced thickness of fur in arid populations
164 compared to mesic, due to the higher temperatures and thus greater need to dissipate heat.
165 We also predicted longer limb length, and smaller overall body size of the individuals in arid
166 regions compared to non-arid individuals, in accordance with Allen's and Bergmann's rules
167 (Alhajeri et al., 2020). Additionally, as mole-rats interact with their surroundings through their
168 teeth, when eating and digging through soil using their incisors, we focus on the skull and the
169 morphological aspects of the head, snout, and teeth. Arid regions will vary from non-arid
170 regions in the soil type and compactness, floral diversity, drier conditions and scarcity of
171 vegetation (Naorem et al., 2023). We predict arid-dwelling individuals to have broader snouts
172 as well as broader zygomatic arches and coronoid processes for attachment of increased
173 muscle mass associated with mastication due to the rocky and compact soil in the arid
174 regions, and xeric plant types found in this habitat.

175 **Methods**

176 **Data collection**

177 Samples of common mole rats were collected from 71 non-breeding individuals across five
178 sites housing discrete populations. Mole-rat individuals were trapped using Hickman live traps

179 (Hickman, 1979) inserted into tunnels located under mounds and were baited with sweet
 180 potato. All sites have been previously documented as having common mole-rats (Spinks et
 181 al., 2000; Visser et al., 2019; Hart et al., 2023). Sites were specifically selected to represent an
 182 aridity gradient, based on an Aridity Index (Table 1). Aridity Index (AI) is a numerical indicator
 183 of the degree of dryness of the climate at a given location (UNEP, 1992). The AI for the study
 184 populations was calculated from climate data (ranging from the years 1981-2020) retrieved
 185 from the ERA5-Land of the European Centre for Medium-Range Weather Forecasts, created
 186 by the Copernicus Climate Change Service (Muñoz-Sabater et al., 2021) with a spatial
 187 resolution of 0.1° by 0.1°. Monthly averaged temperature (T_{air} in °C), total precipitation (t_p in
 188 m), and two-metre dew point temperature ($d2m$ in °C) were used. These combined data were
 189 used to calculate annual aridity index (AI) (equation (1)). Where t_p directly obtained from
 190 ERA5-Land and potential evapotranspiration (PET) calculated from the Romanenko
 191 estimation (equation (2)) (Romanenko, 1961). For equation (2), relative humidity (RH) was
 192 calculated from ERA5-Land $d2m$ (equation (3)).

$$193 \quad AI = \frac{t_p}{PET} \quad (1)$$

$$194 \quad PET = 0.00006 \times (100 - RH) \times (25 + T_{air})^2 \quad (2)$$

$$195 \quad RH = 100 \times 10^{7.591386 \left(\frac{d2m}{d2m + 240.7263} - \frac{T_{air}}{T_{air} + 240.7263} \right)} \quad (3)$$

196

197 Aridity classifications and corresponding AI values, as outlined by UNESCO (1979) and UNEP
 198 (1992) state that where PET is greater than t_p , the climate is considered to be arid (Colantoni
 199 et al., 2015). AI values at each of the five sites used in this study are listed in Table 1 and have
 200 been selected to range in AI across different aridity classifications.

201

202 **Table 1.** List of 5 sites ordered from most arid to least arid, with collection coordinates for
 203 common mole rats at each site, recent Aridity Index values (from the year 2020) for each site
 204 based on climate data taken from ERA5-Land dataset to 2 decimal places. Aridity
 205 classifications correspond to our classifications for the comparison of sites in this study.

Site	Latitude	Longitude	Aridity Index	Aridity Classification
Steinkopf	-29.34531	17.7872	0.04	Arid
No Heep	-30.04253	17.95852	0.07	Arid
Klawer	-31.7013476	18.7446117	0.11	Semi-arid
Darling	-33.406573	18.417538	0.42	Mesic
Somerset West	-34.035613	18.799499	0.86	Mesic

206

207 Steinkopf and No Heep were considered arid, Klawer a semi-arid/intermediate region,
 208 whereas Darling and Somerset West are mesic. The 71 animals used in this study were
 209 maintained in captivity for 22-28 days prior to being euthanised using an overdose of
 210 isoflurane in line with strict veterinary procedures. The Animal Use and Care Committee of
 211 the University of Pretoria evaluated and approved experimental protocols (ethics clearance
 212 No. NAS016/2021) and DAFF section 20 approval (SDAH-Epi-21031811071). Tissues were
 213 harvested by cutting open the abdomen and chest cavity with scissors. Heart, lungs, kidney,
 214 liver, spleen, gastro-intestinal tract, eyes and five pieces of biceps femoris muscle tissue were
 215 collected while blood samples were taken using a syringe and stored in an Eppendorf tube.

216 **Age Class**

217 Age class of individuals was determined by tooth wear of molars and pattern of eruption as
 218 outlined in Bennett et al. (1990). All 71 individuals were non-breeders and comprised a similar
 219 number of males and females in each population subsample used (Table S1).

220 **Organ mass**

221 The organs used in this study were the heart, lungs, kidneys, liver and spleen. Each organ was
 222 weighed upon extraction and stored in various solutions (Supplementary 1) according to the
 223 requirements of the experiments and studies they were subsequently used for. Mass values
 224 were recorded in mg using an electronic precision analytical weighing balance (BIOBASE,
 225 BP1003B), to the nearest 0.01mg for 71 individuals (Steinkopf = 16, No Heep = 19, Klawer =

226 12, Darling = 12 and Somerset West = 12). Dissected bodies were then stored in a -80°C
227 freezer.

228

229 **Body measurements**

230 Measurements were taken on defrosted bodies of 75 individuals (all 71 euthanised
231 individuals, and an additional 4 that died of natural causes; Steinkopf = 16, No Heep = 21,
232 Klauer = 13, Darling = 13 and Somerset West = 12) using 150mm digital callipers (Insize, 1108)
233 to the nearest 0.01mm. The measurements are presented in Table 2 and depicted in Figure
234 S1.

235 **Pelage**

236 Fur absorbance and reflectance were measured using Ocean Optics spectrophotometer
237 (Ocean Optics USB2000, Oxford, UK) measuring between 329nm – 1000nm. Absorbance and
238 reflectance were used to quantify fur colour. Following this, to measure density, each piece
239 of fur was weighed using an electronic precision analytical weighing balance (BIOBASE,
240 BP1003B), to the nearest mg using to measure fur thickness for each of the populations. See
241 Supplementary Information 1 for full methodological details.

242

243 **Table 2.** List of body measurements taken (mm) from 75 common mole rats across five
244 populations varying in Aridity Index along a gradient. Description of each measurement taken
245 of the limbs, paws, teeth and overall body, corresponds to Figure S1 according to number.

	Measurement in mm	Measurement description
1	Body length	Tip of snout to anus
2	Inter-shoulder width	Distance between shoulder blades
3	Inter-hip width	Distance between hips
4	Incisor length	Incisor length top and bottom

5	Fore limb length	Heel of foot to hip/shoulder joint
6	Exposed fore paw length	Front tip of middle toe to fur line
7	Fore paw length	Tip of middle toe to heel
8	Hind limb length	Heel of foot to hip/shoulder joint
9	Exposed hind paw length	Hind tip of middle toe to fur line
10	Hind paw length	Tip of middle toe to heel

246

247 **Skulls**

248 Once body measurements were complete, skulls were extracted. See Supplementary
249 Information 2 for full methodological details.

250 Samples were boxed and shipped on dry ice from University of Pretoria, South Africa, to
251 Queen Mary University of London using World Courier (UK) Limited.

252 **3D imaging**

253 Skull specimens were digitised, and 3D images were created using laser surface scanning
254 (Picza LPX-1200DS 3D Laser Scanner). Circumferential pitch was set to 0.18mm and height-
255 direction pitch was set to 0.10mm. A preliminary scan was carried out in order to determine
256 the resolution. Crania and right lower jaw were scanned for each sample. Left lower jaw for
257 two individuals from each population were also scanned. Both sides of the jaw were assessed
258 in a preliminary analysis on a subsample of the dataset to determine the magnitude of
259 symmetry, which was found to be high (Figure S2), thus it was determined that one-side-only
260 data could be used in the study (Klingenberg, 2002; Cardini, 2016; Cardini, 2017). In total, 14
261 samples were omitted on account of damage to base of the skull and zygomatic arch, leaving
262 57 specimens left in the study (Steinkopf = 10, No Heep = 11, Klawer = 12, Darling = 12 and
263 Somerset West = 12).

264 **3D visualisation**

265 Crania and lower jaws were digitised in 3D using MeshLab, 3D Mesh Processing System
266 Version 2022.02 (Cignoni et al., 2008). A Screened Poisson Surface Reconstruction algorithm
267 (Kazhdan and Hoppe, 2013) was used to build a triangulated mesh out of point cloud data for
268 each specimen, and a PLY file was created for landmarking.

269 **Landmarking**

270 A configuration of 36 3D anatomical landmarks were used which were placed on forms that
271 could be reliably and accurately located and have a clear correspondence between
272 specimens. Focus was placed on the functional parts of the front of the skull and jaw, where
273 the individuals would be interacting with their environment. This consisted of eight landmarks
274 placed around the orbital region, five around the cranial base, four on the supracranium, eight
275 around the upper dentition (incisors and cheek teeth), and ten on the lower jaw (See Table 3
276 and Figure 1). The cranial base is underrepresented due to the placement of the skulls during
277 scanning, so this area has not been scanned clearly enough for landmarking. All landmarking
278 was conducted in Checkpoint Version 2022.12.16.0419 (Stratovan Checkpoint, 2022). Five
279 specimens were landmarked a repeat of five times to ensure intra-observer reliability.

280

281 **Table 3.** Landmarks used for describing cranial and lower jaw shape in common mole rats.
282 Landmark numbers are depicted in Figure 1. Right and left follow standard anatomical
283 directions.

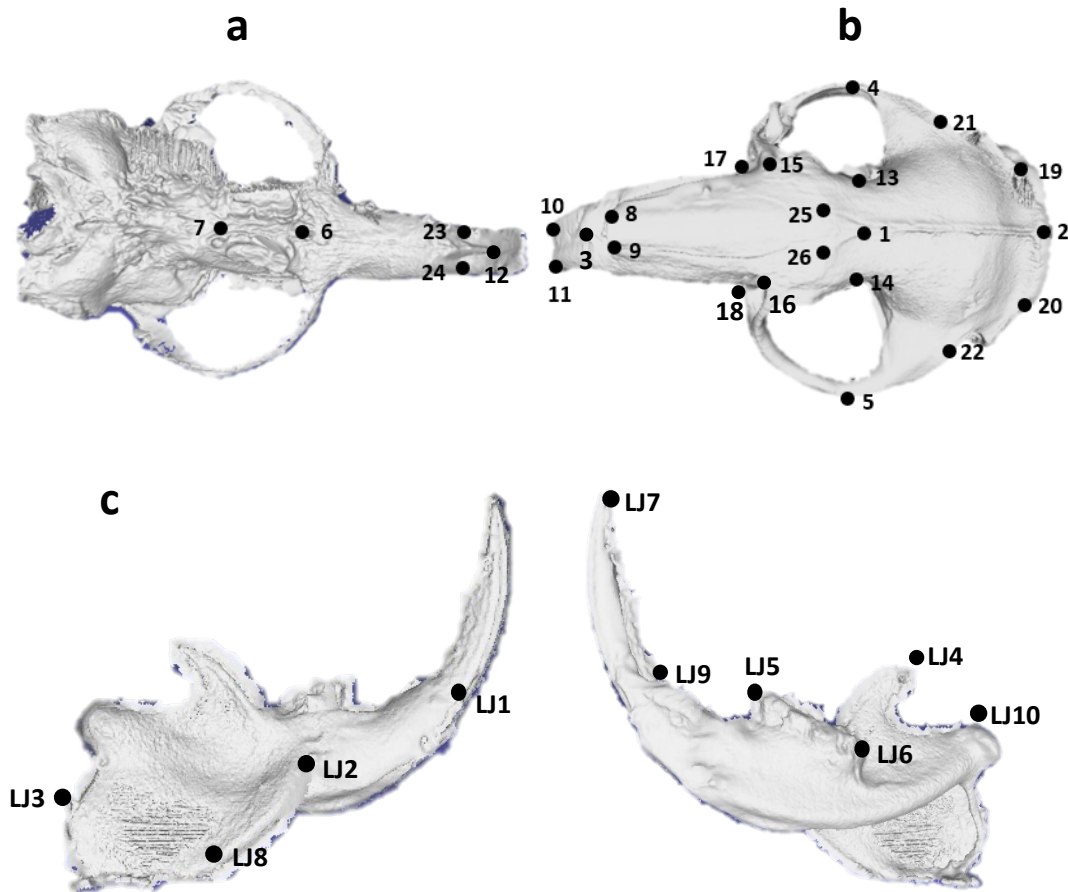
Landmark number	Landmark details
1	Intersection between inter-parietal and inter-frontal sutures
2	Most distal point of the supra-occipital
3	Inter-incisor at the pre-maxillary joint, dorsal side
4	Outer-most point of right zygomatic arch
5	Outer-most point of left zygomatic arch
6	Inter-maxillary suture (Front of top cheek teeth, between left and right cheek teeth rows)
7	Inter-maxillary suture (Back of top cheek teeth, between left and right cheek teeth rows)
8	Intersection between the premaxillary and right nasal suture

9	intersection between the premaxillary and left nasal suture
10	Outer corner of the tip of top right incisor
11	Outer corner of the tip of top left incisor
12	Inter-incisor at the pre-maxillary joint, ventral side
13	Frontal bone, inner point of right temporal fenestrae
14	Frontal bone, inner point of left temporal fenestrae
15	Intersection between the fronto-maxillary suture- right
16	Intersection between the fronto-maxillary suture- left
17	Intersection between the premaxillary-maxillary suture- right
18	Intersection between the premaxillary-maxillary suture- left
19	Lambdoid suture- right
20	Lambdoid suture- left
21	Squamosal-zygomatic joint- right
22	Squamosal-zygomatic joint- left
23	Top incisor at pre-maxillary joint on outer edge- right
24	Top incisor at pre-maxillary joint on outer edge- left
25	Intersection between the parietal and frontal suture- right
26	intersection between the parietal and frontal suture- left
LJ1	Intersection between the incisor-mandible- bottom
LJ2	Intersection between the angular process-mandible joint
LJ3	Back of angular process, placed on the posterior
LJ4	Tip of posterior coronoid process
LJ5	Top of front molar
LJ6	Bottom of back molar at the tooth and mandible joint
LJ7	Tip of incisor
LJ8	Angular process bend
LJ9	Intersection between the incisor and mandible - top
LJ10	Anterior tip of mandibular condyle

284

285

286



287

288 **Figure 1.** Positions of landmarks on cranium and lower jaw (LJ) of 57 *C. h. hottentotus*
 289 specimens. Landmarks are described in Table 3. **A)** Dorsal view of skull, **b)** ventral view of
 290 skull, **c)** medial and lateral view of right lower jaw.

291

292 **Geometric morphometrics**

293 MorphoJ Version 2.0 (Klingenberg, 2011) was used to scale, rotate, and translate the 3D
 294 landmark coordinates to carry out a Procrustes Superimposition (Ross, 2004).

295 **Analyses**

296 All statistical analyses were performed using the statistical software R version 4.2.2 (R Core
 297 Team, 2021). Principal components analyses (PCA) were conducted using the *prcomp*
 298 function and the *factoextra* package in R. PCAs were carried out on the relative mass of organs
 299 (organ mass/body mass) listed under 'Organ mass', to determine variation in the mass of

300 internal organs across 71 individuals and the measurements listed in Table 2, across 75
301 individuals (Table S1).

302 Using the *glm* function, generalised linear models were used to assess population differences
303 for the masses of each organ separately, fur reflectance, fur absorbance and fur thickness.
304 Sex, age class, and population were included as predictor variables, with the addition of body
305 mass for organ mass models. Sex and age class were included in the models to determine if
306 either influence the variation in morphological characteristics. Model selection was
307 determined using the *drop1* function and models with the highest rank were selected using
308 Akaike Information Criterion (AIC) values, thus determining which model best explains our
309 dataset (Grueber et al., 2011). Population and sex remained in the final models for each fur
310 measurement, and sex, age class, body mass and population remained for the organ masses.

311 A PCA was also conducted on the landmarks of the crania and lower jaw separately, for 57
312 individuals to generate a morpho-space of skull shape variation between the populations
313 along an environmental gradient. using the *lda* function from the MASS package in R, a
314 canonical discriminant analysis (CDA) (Williams, 1983) was carried out to support and
315 highlight differences in shape between the populations. This was supported with a pairwise
316 Mahalanobis distance matrix to quantify the distances between group means using
317 multivariate data created using the *mahalanobis* function from the stats package in R. Finally,
318 a pairwise nonparametric multivariate analysis of variance (NPMANOVA) was used using the
319 *pairwise.adonis* and *p.adjust* functions from the pairwiseAdonis R package to test for a
320 significant difference between the Procrustes coordinates, followed by a post-hoc (pairwise)
321 test with a Bonferroni adjustment to find potential differences between populations.

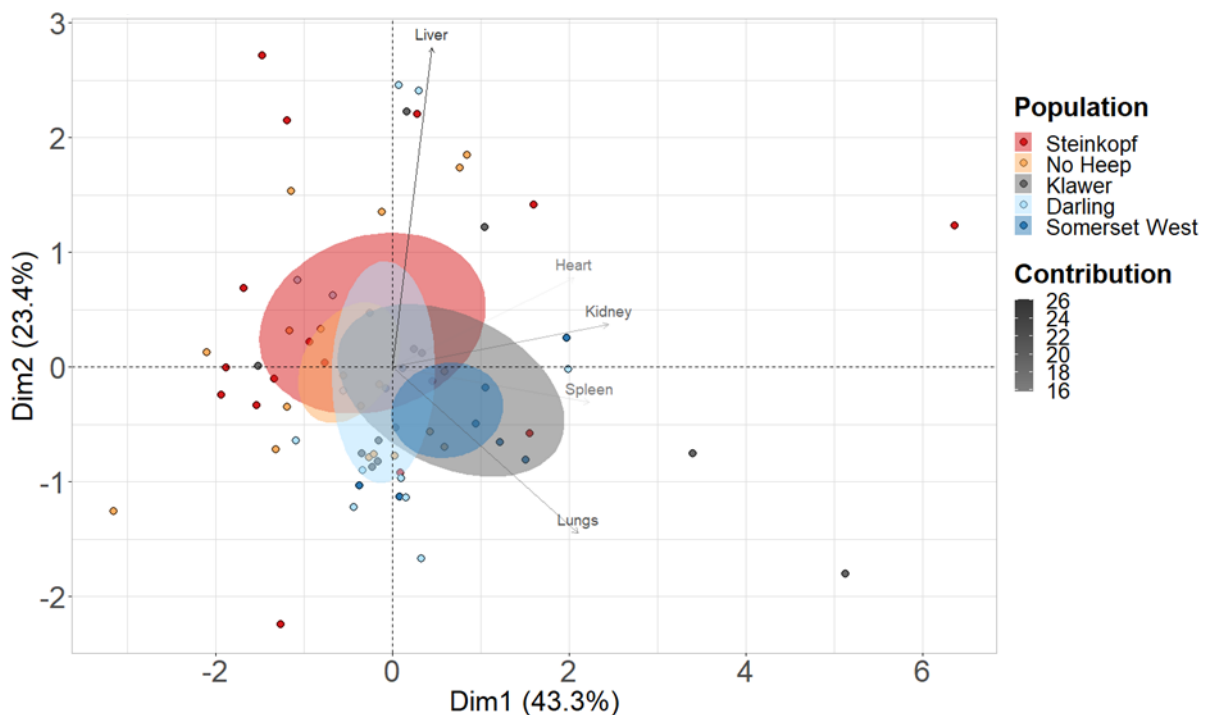
322 Wireframe graphs of the principal component 1 (PC1) and 2 (PC2) from the PCA analysis were
323 generated in MorphoJ (Klingenberg, 2011) to visualise the variation in shape related to the
324 explanatory components which included the highest percentage of shape variation across the
325 populations.

326 **Results**

327 **Organ mass**

328 The PCA of relative organ mass showed that components 1-3 accounted for 81.9% of the
 329 variation in organ mass between populations (Figure 2). PC1 and PC2, displayed 43.3% and
 330 23.4% variance (Table S3), thus, the variation exhibited between populations was well
 331 explained by the variables used in the PCA. The highest contributors were the liver, lungs and
 332 kidneys, showing the majority of differences in relative organ mass (Figure S3). Separation
 333 along the x-axis is shown between the arid and mesic populations in the PCA. Furthermore,
 334 sex was not found to be a discriminating factor and thus variation is not linked to sex (Figure
 335 S4).

336 General linear models (GLMs) for each of the organs demonstrated that body mass was a
 337 significant predictor for each of the organs: heart, (GLM: $F = 8.95$, $df = 9, 59$, $p < 0.001$), lungs
 338 (GLM: $F = 23.41$, $df = 9, 59$, $p < 0.001$), kidneys (GLM: $F = 18.01$, $df = 9, 59$, $p < 0.001$), liver
 339 (GLM: $F = 11.1$, $df = 9, 59$, $p < 0.001$), spleen (GLM: $F = 3.94$, $df = 9, 59$, $p < 0.001$). Spleen mass
 340 was significantly higher in individuals from Somerset West (the least arid population) than all
 341 other populations.



342
 343 **Figure 2.** Principal component analysis based on mass corrected (relative) organ mass of 71
 344 individuals of *C. h. hottentotus* across five populations, Steinkopf, No Heep, Klawer, Darling
 345 and Somerset West. Confidence ellipses are shaded according to population colour and define
 346 the region containing 95% of the samples drawn from the underlying Gaussian distribution.

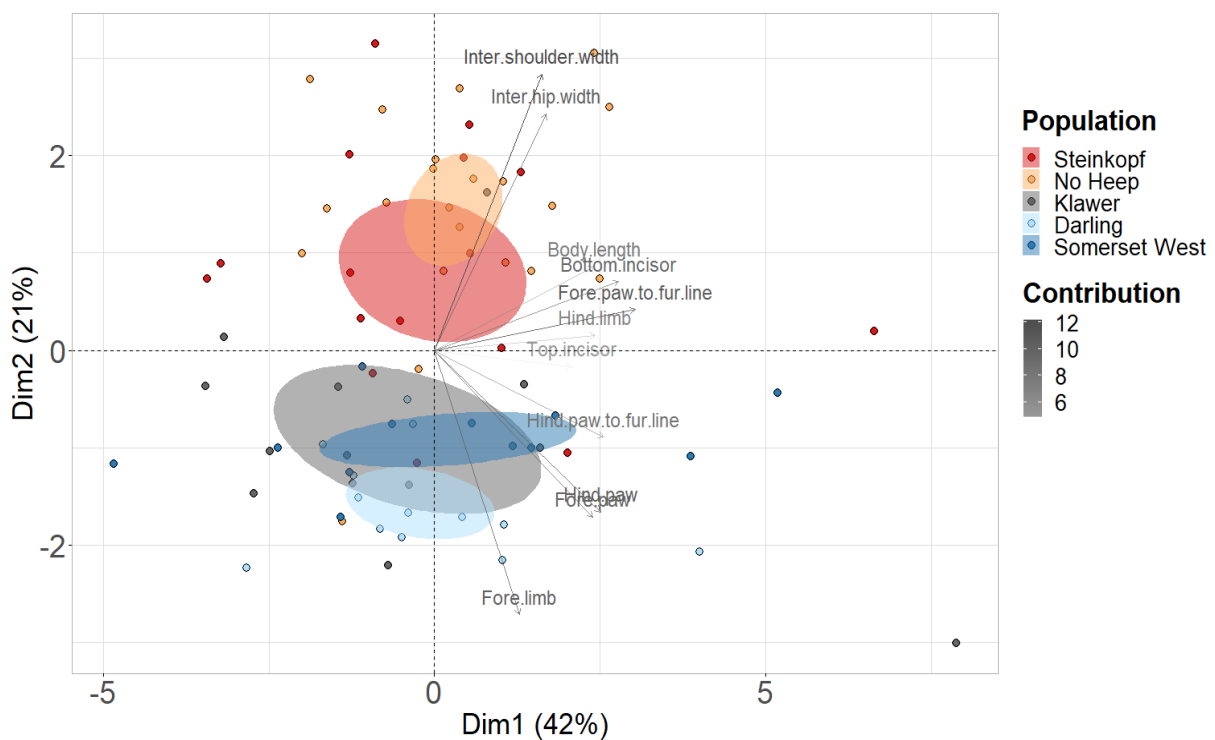
347 The first (Dim1) and second (Dim2) principal components display 43.3% and 23.4% of the total
348 variation, respectively. Contributions of each variable used in the PCA analysis are displayed
349 using a gradient, blue indicating the highest contribution and red the lowest contribution.

350

351 **Body measurements**

352 The principal components (PC) 1-3 accounted for 70.6% of the body and surface skeletal
353 variation. Thus, the differences between populations are well explained by the variables used
354 in the PCA, suggesting that differences between arid and mesic populations are supported by
355 the measurements used. PC1 and PC2 displayed 42% and 21% variance (Table S4). The highest
356 contributors were fore limb length and inter-shoulder width; the majority of shape variation
357 between populations involves changes in limb length and shoulder span (Figure S5).
358 Separation along the y-axis is shown between the arid and mesic populations in the PCA
359 (Figure 3). Additionally, sex was not found to be a discriminating factor and, thus, variation is
360 not linked to sex (Figure S6).

361



362

363 **Figure 3.** Principal component analysis based on surface skeletal body measures of 75
364 individuals of *C. h. hottentotus* across five populations, Steinkopf, No Heep, Klawer, Darling
365 and Somerset West. Confidence ellipses are shaded in according to population colour and
366 define the region containing 95% of the samples drawn from the underlying Gaussian
367 distribution. The first and second principal component display 42% and 21% of the total
368 variation, respectively. Contributions of each variable used in the PCA analysis are displayed
369 using a gradient, blue indicating highest contribution and red lowest contribution.

370

371 **Pelage**

372 No significant difference was found between the populations in fur reflectance (GLM, $F=2.06$,
373 $p=0.1$) and absorbance (GLM, $F=2.13$, $p=0.09$). Additionally, no difference in fur thickness was
374 found between the populations (GLM, $F= 1.7502$, $p=0.17$).

375

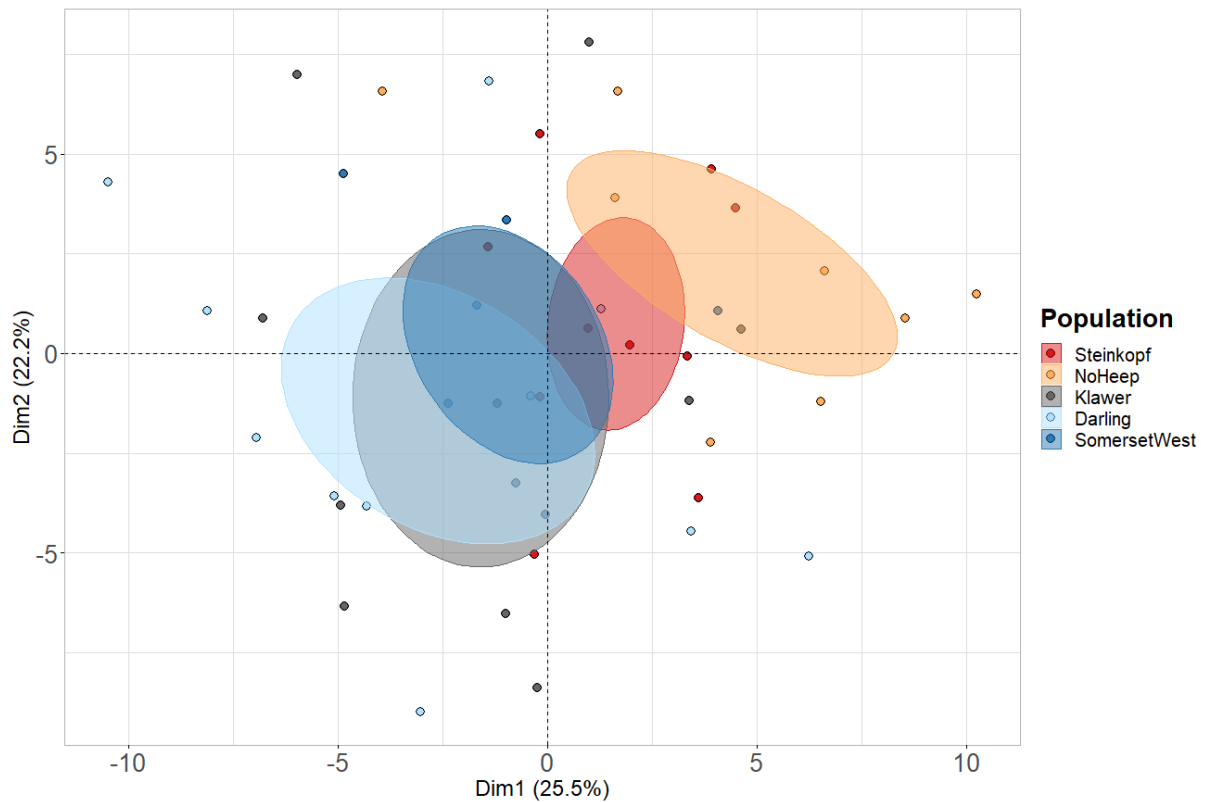
376 **Morphometrics**

377 **PCA**

378 Principal components (PC) 1-3 accounted for 65.23% and 65.97% of the variation in the crania
379 and lower jaw respectively, thus, the variables used in the PCA can moderately explain the
380 observed differences between populations (Figure 5). PC1 and PC2 displayed 25.5% and
381 22.2% variance for the crania (Table S5) and 25.1% and 21.1% in the lower jaw (Table S6); the
382 majority of shape variation involves changes in the depth of the skull and shape of the
383 zygomatic arches (Landmarks 1, 13/14, 21/22 and 25/26). PC1 demonstrates separation
384 between the mesic and arid populations with high intra-group variation. Additionally, sex was
385 not found to be a discriminating factor and thus variation is not linked to sex (Figure S7).

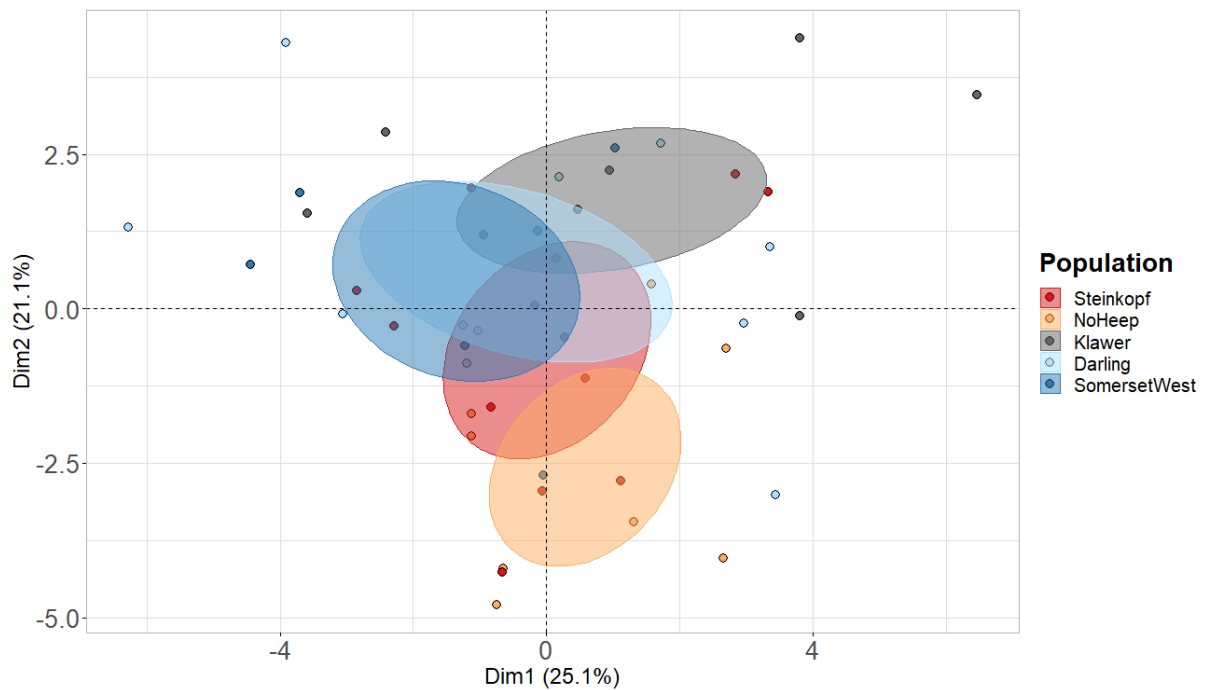
386

a



387

b



388

389 **Figure 4.** Principal component analysis based on landmark data of 57 individuals of *C. h.*
 390 *hottentotus* across five populations, Steinkopf, No Heep, Klawer, Darling and Somerset West
 391 for **a)** crania and **b)** lower jaw. Confidence ellipses are shaded in according to population
 392 colour and define the region containing 95% of the samples drawn from the underlying

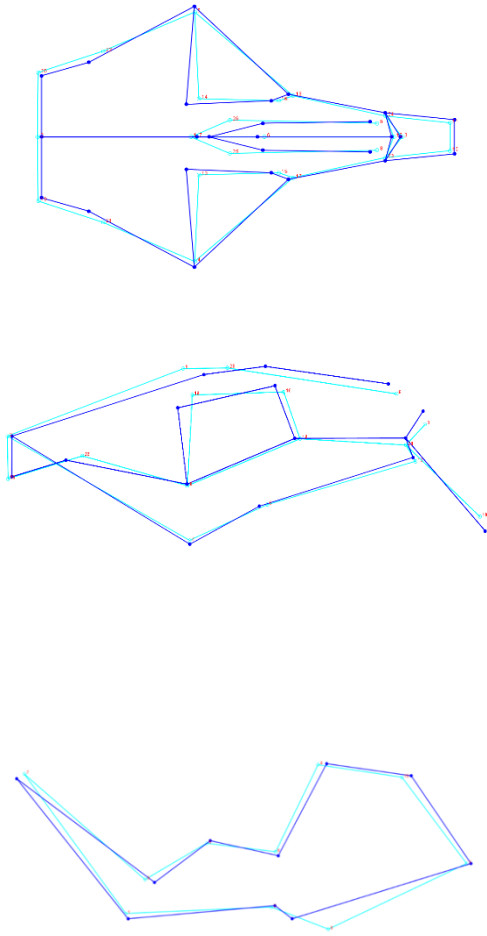
393 Gaussian distribution. The first and second principal component display 25.5% and 22.1% of
394 the total variation, respectively for the crania, and 25.1% and 21.1% for the lower jaw.

395

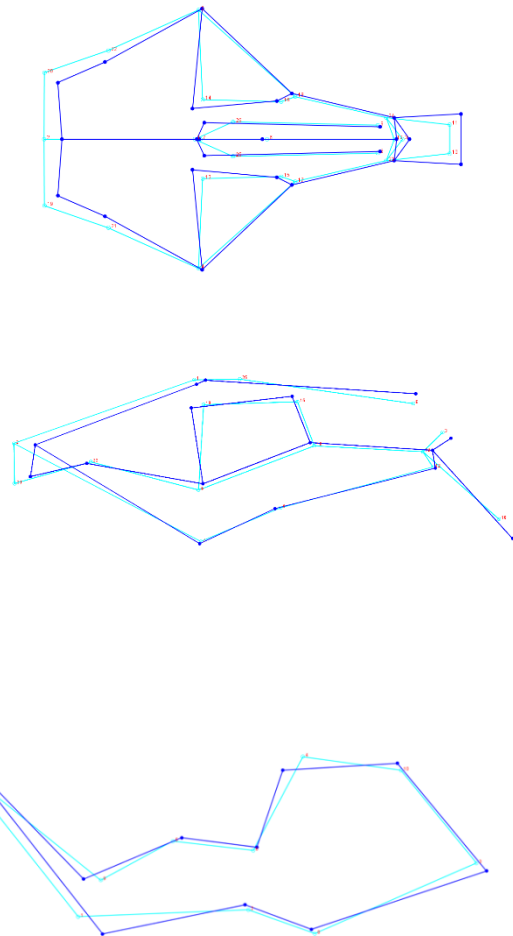
396 Landmarks in PC1 deviated the most at landmarks 8 (LJ) and 1, 13/14, 21/22 and 25/26
397 (crania). These are where landmarks outline the zygomatic arches, dorsal side of the skull and
398 angular process of the lower jaw. These deviations increased in size along the x-axis of the
399 PCA plot, indicating that more arid populations of the mole-rats have greater dorsoventral
400 depth of skulls and angular processes of the lower jaw that are more dorsally expanded
401 (Figure 4). PC2 shows similar variation to PC1 in landmarks 1, 13/14, 21/22 and 25/26 (crania)
402 indicating greater procumbence in the upper incisors. PC2 also showed an increase in size of
403 landmarks 10/11 (crania), indicating that deviations increased in size along the axis.
404 Additionally, there was a decrease in the distance between landmarks LJ1 and LJ9 of the lower
405 jaw, increasing the depth of the mandible. Landmark LJ4 decreases in height (Figure 5).

406

a



b



407

408 **Figure 5.** Wireframe graphs for **a)** PC1 and **b)** PC2 showing the average shape transformation
409 of the crania and lower jaw (LJ) along dimension 1 from left (dark blue) to right (light blue) on
410 the x-axis of the PCA plot for PC1, and along dimension 2 from bottom to top on the y-axis for
411 PC2 (See Figure 5a).

412

413 **Canonical discriminant analysis (linear discriminant analysis)**

414 Canonical discriminant analysis (CDA) showed that arid and mesic populations cluster
415 separately (Figure 6). The pairwise square Mahalanobis distance and probability values reveal
416 that No Heep shows the greatest difference to all other populations (Table 4). Darling and
417 Somerset West, the two least arid populations, are the most similar populations. Darling and

418 Klawer, and Somerset West and Klawer have low values, indicating that there are similarities
 419 between these populations as well (Figure S8 and Figure S9).

420

421 **Table 4.** Pairwise square Mahalanobis distance matrix among five populations of *C. h.*
 422 *hottentotus*. Pairwise distances are calculated from 36 landmarks across the crania (1) and
 423 lower jaw (2) of 57 specimens. All values are $\times 10^6$.

1	Steinkopf	No Heep	Klawer	Darling	Somerset West
Steinkopf	0	12.42	14.83	14.23	28.13
No Heep	12.42	0	31.53	20.83	30.85
Klawer	14.83	31.53	0	7.26	11.59
Darling	14.23	20.83	7.26	0	9.86
Somerset West	28.13	30.85	11.59	9.86	0

424

2	Steinkopf	No Heep	Klawer	Darling	Somerset West
Steinkopf	0	22.43	5.34	6.36	6.28
No Heep	22.43	0	6.71	8.71	13
Klawer	5.34	6.71	0	2.64	3.51
Darling	6.36	8.71	2.64	0	2.73
Somerset West	6.28	13	3.51	2.73	0

425

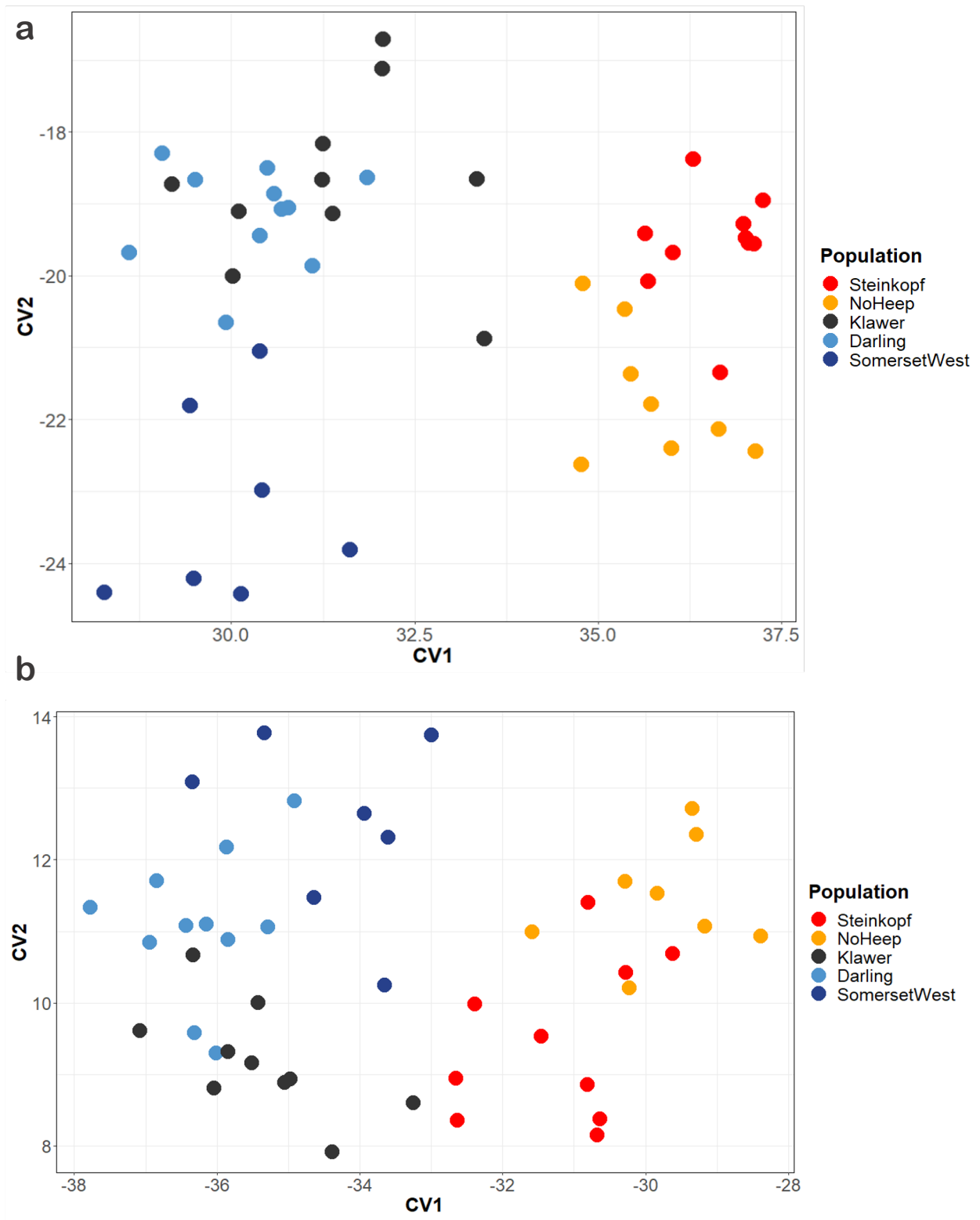
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431

432 **Figure 6.** Distribution of individuals as explained by the first two canonical variates (CV1 and
 433 CV2) derived from landmark data of 57 individuals of *C. h. hottentotus* across five populations,
 434 Steinkopf, No Heep, Klawer, Darling and Somerset West for **a)** the crania and **b)** the lower
 435 jaw.

436 **Non-parametric pairwise MANOVA**

437 A non-parametric pairwise MANOVA and a post-hoc test with a Bonferroni adjustment
 438 showed that there were no significant differences between any of the populations
 439 (NPMANOVA: $F = 1.04$, $df = 4$, $p = 0.4176$) for the crania and a significant difference between
 440 No Heep and Klawer only (NPMANOVA: $F = 1.51$, $df = 4$, $p = 0.124$) for the lower jaws (Table
 441 5).

442

443 **Table 5.** Results of a post-hoc Bonferroni adjustment of landmark data from 57 individuals of
 444 *C. h. hottentotus* populations from five populations for the crania (1) and lower jaw (2).

1	df	Sum of squares	F Model	R²	p value
Steinkopf vs No Heep	1	5.04	0.71	0.043	0.58
Steinkopf vs Klawer	1	4.42	0.72	0.038	0.59
Steinkopf vs Darling	1	13.03	1.66	0.08	0.14
Steinkopf vs Somerset West	1	3.34	0.73	0.046	0.58
No Heep vs Klawer	1	13.39	1.53	0.088	0.19
No Heep vs Darling	1	15.42	1.47	0.079	0.22
No Heep vs Somerset West	1	9.14	1.22	0.086	0.29
Klawer vs Darling	1	7.66	0.83	0.042	0.49
Klawer vs Somerset West	1	1.41	0.22	0.015	0.94
Darling vs Somerset West	1	5.6	0.67	0.04	0.63

2	df	Sum of squares	F Model	R²	p value
Steinkopf vs No Heep	1	5.096	1.54	0.088	0.2
Steinkopf vs Klawer	1	5.77	1.33	0.069	0.25
Steinkopf vs Darling	1	1.19	0.29	0.015	0.88
Steinkopf vs Somerset West	1	2.06	0.64	0.041	0.62
No Heep vs Klawer	1	11.74	3.05	0.16	0.03
No Heep vs Darling	1	5.71	1.57	0.084	0.17
No Heep vs Somerset West	1	7.15	2.94	0.18	0.05
Klawer vs Darling	1	5.41	1.18	0.058	0.29
Klawer vs Somerset West	1	10.72	2.84	0.16	0.04
Darling vs Somerset West	1	2.68	0.75	0.045	0.49

445

446

447

448 **Discussion**

449 Variation in morphology and anatomy was found between populations of the common mole-
450 rats along an aridity gradient. The shape of the skull and body, showed variation along the AI
451 that represents the environments where the populations were collected. Overall, a greater
452 mass of the liver was found in arid populations of *C. h. hottentotus*, compared with those from
453 the intermediate and mesic regions. Variations in body measurements were found between
454 the arid populations and the intermediate and mesic, with arid populations having individuals
455 with shorter forelimb lengths and larger inter-shoulder widths. No significant differences in
456 fur colour and thickness were observed between these populations. Finally, geometric
457 morphometric analyses demonstrated variation in skull shape between mesic and arid
458 populations. Arid populations show evidence of greater depth of skull, and shape variation in
459 features such as angular processes of the lower jaw, and zygomatic arches, suggesting the
460 attachment of larger masticatory muscles.

461 **Organ mass**

462 A larger liver and kidney mass was observed in the specimens derived from arid regions. The
463 roles of the kidney include water regulation and filtration, and detoxification of substances
464 absorbed by the digestive system (Brzoska et al., 2003). Greater kidney mass could indicate
465 links to physiological responses to heat and dehydration such as increased water regulation
466 and water turnover, as well as a greater ability for water retention which can be utilised
467 during periods of stress, starvation, and dehydration (Jacobs et al., 2020; Jacobs et al., 2022).
468 Other rodent and mole-rat studies have explored the differences in kidney mass, size,
469 function, and metabolism in arid and mesic regions, with similar findings (Al-Kahtani et al.,
470 2004; Jacobs et al., 2022; Jackson et al., 2004). Many bulbs found in the Northern Cape, where
471 the arid populations in this study occur, have been found to be toxic, such as the bulbs of the
472 *Drimia* plants (Manganyi et al., 2021). *Drimia* bulbs contain high levels of cardiac glycosides
473 and are widespread across the Northern Cape (Bozorgi et al., 2017), thus, the liver and
474 potentially the kidneys of individuals in these arid regions may be playing a role in
475 detoxification of these food items.

476 A large proportion (N=7 out of 12) of individuals from Somerset West showed enlarged
477 spleens, and this could be linked to immunity as Somerset West is home to two other
478 sympatric mole-rat species, *Georychus capensis* (Cape mole-rats) and *Bathyergus suillus* (Cape

479 dune mole-rats), (Robb et al., 2016; Thomas et al., 2013). The spleen is the primary producer
480 of blood cells in the mammalian body, and has important functions involved in
481 haematopoiesis (formation of blood cells such as red blood cells, macrophages and
482 antibodies), blood filtration and immunity (Emmrich et al., 2019). An enlarged spleen is often
483 associated with an immune response to infection or inflammation (Cheng et al., 2017).
484 However, Cheng et al. (2017) observed that naked mole-rats (*H. glaber*) have enlarged spleens
485 for their size, relative to lab mice, and is believed to combat the risk of infection and disease
486 transmission in large and tight-knit colonies containing many individuals (Bégay et al., 2022).
487 The anatomy of the spleen in relation to colony rank in naked mole-rats has also been
488 investigated whereby, enlarged spleens were shown in individuals with a higher rank within
489 the colony (Bégay et al., 2022). Variation in spleen size has been proposed to be regulated by
490 social interactions, and to provide immunological advantages to higher ranking individuals
491 that patrol the colony. These individuals are also often involved in colony defence and, thus,
492 have an increased risk of contact with predators and intruders carrying unfamiliar pathogens
493 (Bégay et al., 2022). The individuals with enlarged spleens may be from colonies that share
494 territories with other heterospecific species. The interaction with individuals outside of the
495 colony may lead to exposure to foreign microorganisms, leading to enlarged spleens in
496 individuals involved in patrol or colony protection roles. It is also possible that, by chance,
497 individuals from higher ranks were sampled from Somerset West, compared to the arid
498 populations, as per observations from Bégay et al. (2022).

499 Body mass variation was a significant influencer of organ mass, and arid dwelling individuals
500 had a larger body mass relative to the mesic dwelling individuals. A greater body mass in arid
501 biomes could be due to the positive relationship between size and water conservation ability
502 (Naya et al., 2017). Arid populations are likely under selection pressures that require greater
503 water retention to avoid dehydration, thus increased body mass. General linear models
504 showed body mass as the only significant predictor of organ mass for all organs except the
505 spleen.

506 **Body and skull morphology**

507 The PCA for the 10 body measurements of *C. h. hottentotus* showed differences between the
508 arid and mesic populations; the highest contributing variables were forelimb length and inter-

509 shoulder width. Converse to Allen's rule, forelimbs were shorter in mole-rats from arid
510 populations compared to those from the mesic populations. Limb length is likely influenced
511 by alternative selective pressures than those presented purely by climatic conditions, but
512 further investigation into limb function and usage is needed before firm conclusions can be
513 drawn. There is, perhaps, a trade-off between thermoregulatory pressures of arid
514 environments and selection for alternative traits such as locomotion and mobility in burrow
515 systems. Several studies have shown similar patterns in limb length varying conversely to
516 Allen's rule, across the rodent Order (Alhajeri et al., 2020), and in other subterranean and
517 semi-fossorial rodents (Bidau et al., 2011; Lindsay, 1987). Longer limb length is predicted in
518 warmer climates to increase distance from the hot substrate; an adaptation that is redundant
519 in subterranean species. Additionally, highly adaptive features are required for mobility in
520 burrows or shelters, thus fossoriality may be a stronger influencer on limb length than aridity.
521 In the mesic regions, both *Bathyergus suillus* and *B. janetta* species are found sympatrically
522 with common mole-rats, both of these species utilise claw digging as well, to facilitate digging
523 with their teeth (Bennett and Faulkes, 2000). Soft, sandy soils are found in the mesic regions
524 and as such, burrows will collapse when digging, so *B. suillus* and *B. janetta* need the ability
525 to move soil out of the way with their feet as they dig. The longer limb length found in the
526 mesic populations suggest that these populations could be tending towards utilising some
527 degree of forelimb movement to facilitate digging in soft soils. Indeed, front paw length was
528 the fourth highest contributor. Montoya-Sanhueza et al. (2022) found that solitary species of
529 African mole-rats exhibit increased limb bone specialisation compared to social species, due
530 to the shared costs of digging in colonial species. This may further suggest that arid
531 populations of common mole-rats are living in larger colonies, and that digging roles are
532 shared between a larger number of individuals.

533 Cranial and lower jaw shape varies along the environmental gradient and shows evidence of
534 separate morphotypes in arid and mesic habitats (Barčiová et al., 2009). Arid populations
535 demonstrated skull shape changes linked to increased muscle attachment, such as broader
536 rostra, wider zygomatic arches, larger temporal fossae and greater depth of the skull.
537 Furthermore, the arid individuals also had a larger inter-shoulder width than mesic
538 individuals. The harder soil found in arid regions necessitates stronger muscles to loosen and
539 remove the soil (Kraus et al., 2022). The difference in inter-shoulder width may be linked to

540 neck and chest muscle attachments of the different populations and the increased muscle
541 mass of the skull may require greater muscular and skeletal anatomy of the skull and
542 shoulders for support. The greater inter-shoulder width of individuals in arid populations also
543 suggests increased muscle mass potentially needed for the forelimbs to aid in movement
544 through harder-packed soil. The AFDH suggests increased demand for foraging due to sparse
545 geophyte distribution in arid populations of common mole-rats which may require increased
546 digging compared to the mesic populations to find food, therefore requiring larger muscles
547 for this constant foraging. Evidence exists for more extensive tunnel systems in the arid
548 populations compared to the mesic populations (Spinks et al., 2000), and this may necessitate
549 larger skull features that support masticatory muscle attachments, for the greater muscular
550 mass. This suggests increased muscle mass indicates increased strength to dig and extend the
551 tunnel systems. Mammalian masticatory morphology is known to be a highly plastic region
552 of the skull. The functional morphology of crania and masticatory musculature in several
553 mammalian species has been studied in relation to ecological factors such as diet, habitat,
554 locomotory and activity patterns and found to be linked to diet and bite force (Gomes
555 Rodrigues et al., 2016; Gomes Rodrigues and Damette, 2023). This indicates that larger muscle
556 masses are related to greater bite force and increased chewing, particularly regarding
557 dentition and the lower jaw (Becerra et al, 2011; Borges et al., 2017; Gomes Rodrigues et al.,
558 2023). Hystricognath rodents have shown variations in morphology associated with
559 mastication in line with variations in habitat and diet (Hautier et al., 2012). Similar patterns
560 have been found in other species such as punaré rats (*Thrichomys apereoides*) (Monteiro et
561 al., 2003), where individuals were sampled along an environmental gradient and populations
562 from arid environments had larger coronoid processes, larger jugals, and wider snouts
563 compared to non-arid populations. This is thought to be related to bite force and linked to
564 the vegetation type of the region (Monteiro et al., 2003). As such, bite force has a strong
565 influence on muscle development, even within a species' lifetime. Evidence is emerging that
566 bite force may also have wider implications regarding social structure and reproductive
567 success, and thus increased selection pressure (Kraus et al., 2022). The greater skull depth
568 observed in the arid populations suggests a greater action of the chisel digging method, likely
569 required due to the hardness of the soil (McIntosh and Cox, 2016a). Furthermore, the
570 different biomes of the arid and mesic regions consist of different vegetation (Wright and
571 Samways, 1996). Arid populations of common mole-rats will be feeding on desert geophytes

572 that are tougher, and with thicker skins to reduce water loss in arid environments (Robb et
573 al., 2016), and thus will require more chewing to break down. This increased need for stronger
574 masticatory action in arid regions could also explain the larger inter-shoulder widths and
575 larger attachments for maxillary musculature in these populations. The clustering of the
576 intermediate population for both body and skull measurements shows that there are more
577 similarities in the diet and soil types of Klawer to the mesic population than to that of the arid.
578 Future work could explore the mass of masticatory muscles, such as those of the masseteric
579 complex and temporal region. This would allow for comparison of the relative muscle mass
580 between the arid and mesic populations in relation to the use of incisors. Magnetic Resonance
581 Microscopy may enable imaging and quantification of muscle volume, without the need for
582 excision (Driehuys et al., 2008). Paired with data on soil hardness, this study would shed light
583 on the direct effects of soil type on skull morphology of common mole-rats.

584

585 **Conclusion**

586 Aridity embraces a series of habitats that covers a great deal of the Earth's surface and, thus,
587 it is important to understand how environmental factors directly influence morphological
588 traits across populations of the same species, along an environmental gradient. We have
589 found evidence for extensive variation in body and skull morphology of populations of *C. h.*
590 *hottentotus* found in arid and mesic populations. The separate clustering of mesic and arid
591 individuals with different morphologies suggests the environment does play a significant role
592 in the morphological variation seen in populations of *C. h. hottentotus*. These differences are
593 counter to what is expected in mammals and indicate arid and mesic mole-rat populations
594 show unique adaptations in response to the food source variation and soil composition of the
595 different biomes. This study contributes to a better understanding of the intra-specific
596 morphology of mole-rats distributed along an aridity gradient, and skull morphology as an
597 adaptive response to aridity and food distribution. We can also gain an insight into how
598 subterranean species can change their highly specified subterranean morphology to cope
599 with climatic differences. This can help to infer the level of adaptive morphological specificity
600 within species and highlight the challenges they may face in changing climates and whether

601 it is possible for a species to change its morphology, and rapidly enough, to cope with climate
602 changes.

603

604 **ETHICS**

605 The Animal Use and Care Committee of the University of Pretoria evaluated and approved
606 experimental protocol (ethics clearance No. NAS016/2021) and DAFF section 20 approval
607 (SDAH-Epi-21031811071). Relevant provincial animal capture permits were obtained
608 (Western Cape: CN44-87-13780; Northern Cape: FAUNA 0419/2021, FAUNA 042/2021;
609 Gauteng: CPF6-0124). All methods were performed following the relevant guidelines and
610 regulations. In addition, all experimental procedures were carried out under the
611 recommendations in the Guide for the Care and Use of Laboratory Animals of the National
612 Institutes of Health.

613

614 **CONFLICT OF INTEREST**

615 We have no competing interests.

616

617 **DATA ACCESSABILITY**

618 All data are available as an electronical supplementary file.

619

620 **AUTHORS' CONTRIBUTIONS**

621 Conceptualisation, methodology and planning, H.N.M, S.J.P, C.G.F, D.W.H. and N.C.B. Data
622 collected by H.N.M, D.W.H., N.C.B., A.K.J.v.V. Access to equipment provided by J.B.
623 Methodological development, H.N.M, S.J.P, C.G.F, D.W.H. and N.C.B. Data analysis was
624 performed by H.N.M. All authors contributed to the intellectual interpretation of the results.
625 The initial draft of the paper was written by H.N.M. with contribution and edits by all authors.

626

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