

1 **Running head:** CRANIAL MORPHOLOGICAL DIVERSITY IN  
2 TENRECS

3 Morphological diversity of tenrec  
4 (Afrosoricida, Tenrecidae) crania is greater  
5 than their closest relatives, the golden  
6 moles (Afrosoricida, Chrysochloridae)

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14 diversity, tenrecs

## <sup>15</sup> **Abstract**

## 16 Introduction

17 Morphological diversity has long attracted the attention of biologists.  
18 There are many famous examples of morphological diversity including  
19 beak morphologies in Darwin's finches, body and limb morphologies in  
20 Caribbean *Anolis* lizards and pharyngeal jaw diversity in cichlid fish  
21 (Gavrillets & Losos, 2009). Apart from a few examples (REFS), it is  
22 common to study morphological diversity from a qualitative rather than  
23 quantitative perspective (REFS). However, it is important to quantify  
24 morphological diversity because it has implications for studies of adaptive  
25 radiations (Losos, 2010), convergent evolution (REF) and our  
26 understanding of biodiversity (Roy & Foote, 1997).

27 Tenrecs are an example of a morphologically diverse group  
28 (Soarimalala & Goodman, 2011; Olson & Goodman, 2003). The Family  
29 contains 34 species, 31 of which are endemic to Madagascar (Olson, 2013).  
30 Body sizes of tenrecs span three orders of magnitude (2.5 to > 2,000g)  
31 which is a greater range than all other Families, and most Orders, of  
32 living mammals (Olson & Goodman, 2003). Within this vast size range  
33 there are tenrecs which convergently resemble shrews (*Microgale* tenrecs),  
34 moles (*Oryzorictes* tenrecs) and hedgehogs (*Echinops* and *Setifer* tenrecs)  
35 (Eisenberg & Gould, 1969) even though they are not closely related to  
36 these species (Stanhope et al., 1998). However, morphological diversity in  
37 tenrecs has not been quantified.

38 Morphological diversity is difficult to quantify. Studies are inevitably  
39 constrained to measure the diversity of specific traits rather than overall  
40 morphologies (Roy & Foote, 1997). Different trait axes (such as cranial  
41 compared to limb morphologies) may yield different patterns of

42 morphological diversity (REF) Furthermore, linear measurements of  
43 morphological traits can restrict our understanding of overall  
44 morphological variation (REF). However, geometric morphometric  
45 approaches (Rohlf & Marcus, 1993; Adams et al., 2013) provide more  
46 detailed insights into morphological variation.

47 Here we present the first quantitative investigation of morphological  
48 diversity in tenrecs. We use geometric morphometrics to compare cranial  
49 morphological diversity in tenrecs to their sister taxa, the golden moles  
50 (Afrosoricida, Chrysochloridae). Tenrecs inhabit a wider variety of  
51 ecological niches (Soarimalala & Goodman, 2011) than golden moles  
52 (Bronner, 1995) so we expected tenrecs to be more morphologically  
53 diverse than their closest relatives. However, we find no significant  
54 difference in the diversity of cranial morphologies between the two  
55 groups. It is only when we restricted our data to include a subsample of  
56 the morphologically similar *Microgale* tenrec Genus that we found tenrecs  
57 to be more morphologically diverse than golden moles. Our results  
58 demonstrate the importance of using quantitative methods to assess  
59 otherwise subjective estimates of morphological diversity. We show that  
60 the apparently high morphological diversity in tenrecs is not necessarily  
61 reflected in all morphological traits.

## 62 **Materials and Methods**

### 63 **Morphological data collection**

64 One of us (SF) photographed cranial specimens of tenrecs and golden  
65 moles at the Natural History Museum London (BMNH), the Smithsonian

66 Institute Natural History Museum (SI), the American Museum of Natural  
67 History (AMNH), Harvard's Museum of Comparative Zoology (MCZ)  
68 and the Field Museum of Natural History, Chicago (FMNH). We  
69 photographed the specimens with a Canon EOS 650D camera fitted with  
70 an EF 100mm f/2.8 Macro USM lens using a standardised procedure to  
71 minimise potential error (see supplementary material for details).

72 We collected pictures of the skulls in dorsal, ventral and lateral views  
73 (right side of the skull). A full list of museum accession numbers and  
74 details on how to access the images can be found in the supplementary  
75 material.

76 In total we collected pictures from 182 skulls in dorsal view (148  
77 tenrecs and 34 golden moles), 173 skulls in ventral view (141 tenrecs and  
78 32 golden moles) and 171 skulls in lateral view (140 tenrecs and 31 golden  
79 moles) representing 31 species of tenrec (out of the total 34 in the family)  
80 and 12 species of golden moles (out of a total of 21 in the family (Asher  
81 et al., 2010)). We used the taxonomy of Wilson and Reeder (2005)  
82 supplemented with more recent sources (Olson, 2013) to identify our  
83 specimens.

84 We used a combination of both landmarks (type 2 and type 3,  
85 (Zelditch et al., 2012)) and semilandmarks to characterise the shapes of  
86 our specimens. Figure 1 shows our landmarks (points) and  
87 semilandmarks (outline curves) for the skulls in each of the three views.  
88 Corresponding definitions of each of the landmarks can be found in the  
89 supplementary material.

90 We digitised all landmarks and semilandmarks in tpsDIG, version 2.17  
91 (Rohlf, 2013). We re-sampled the outlines to the minimum number of

evenly spaced semilandmark points required to represent each outline accurately (MacLeod, 2013, details in supplementary material). We used TPSUtil (Rohlf, 2012) to create "sliders" files (Zelditch et al., 2012) that defined which points in our tps files should be treated as semilandmarks. We conducted all subsequent analyses in R version 3.0.2 (R Core Team, 2014) within the geomorph package (Adams et al., 2013). We used the gpagen function to run a general Procrustes alignment (Rohlf & Marcus, 1993) of the landmark coordinates while sliding the semilandmarks by minimising Procrustes distance (Bookstein, 1997). We used these Procrustes-aligned coordinates of all species to calculate average shape values for each species ( $n = 43$ ) which we then used for a principal components analysis (PCA) with the plotTangentSpace function (Adams et al., 2013).

## Calculating morphological diversity

## Results

### Morphological disparity in tenrecs and golden moles

Figure 2 depicts the morphospace plots derived from our principal components analyses of average Procrustes-superimposed shape coordinates for each species in our skull and mandible data respectively. We used the principal components axes which accounted for 95% of the cumulative variation (number of axes:  $n = 7$  (dorsal),  $n = 8$  (ventral),  $n = 8$  (lateral) and  $n = 12$  (mandibles)) to calculate the disparity of each Family.

Tenrecs and golden moles clearly have very different cranial and

115 mandible morphologies: in each analysis, the families occupy significantly  
116 different areas of morphospace (npMANOVA, table 3). In our analyses of  
117 the three different views of the skulls, there is an overall trend for tenrecs  
118 to have higher disparity than golden moles. However, none of these  
119 differences are statistically significant (table 1).

120     There is a less clear pattern from our analysis of disparity in  
121 mandibles. Two of our four metrics indicate that golden moles have  
122 significantly higher disparity in the shape of their mandibles than tenrecs  
123 (table 1) although one metric (sum of ranges) gives the opposite result.

124     The three curves at the back of the mandibles (figure ??) place a  
125 particular emphasis on shape variation in the posterior of the bone; the  
126 ramus, coronoid, condylar and angular processes. Therefore, higher  
127 disparity in golden mole mandibles compared to tenrecs could be driven  
128 by greater morphological variation in these structures. To test this idea,  
129 we repeated our morphometric analyses of the mandibles with a reduced  
130 dataset of points; just the seven landmark points and one single curve at  
131 the base of the jaw between landmarks 1 and 7 (figure ??). When we  
132 compared disparity with this reduced data set we found that golden  
133 moles no longer had significantly higher disparity than tenrecs (table 1).

## 134 **Morphological disparity in non-*Microgale* tenrecs and** 135 **golden moles**

136 We repeated our disparity comparisons with a subset of the tenrec  
137 specimens to remove the large and phenotypically similar *Microgale* tenrec  
138 Genus. In this case we found that tenrecs have significantly higher  
139 disparity than golden moles when the skulls are analysed in lateral view

140 (table 2). However, none of the other comparisons in any of the analyses  
141 were significant.

## 142 Discussion

143 Our analyses are the first quantitative investigation of morphological  
144 disparity in tenrecs. We show that tenrecs' cranial morphologies are no  
145 more diverse than their closest relatives and therefore phenotypic variety  
146 in tenrecs is perhaps not as exceptional as it first appears.

147 When we compared the diversity of skull shapes in the two Families,  
148 we found a trend towards higher disparity in tenrecs compared to golden  
149 moles but none of these differences were significant (table 1). Even when  
150 we removed the phenotypically similar *Microgale* Genus, tenrecs were still  
151 no more diverse than golden moles in most of the analyses of their skull  
152 shapes (table 2).

153 In contrast to these results for the skulls, two of our disparity metrics  
154 indicate that golden moles have more disparate mandible shapes than  
155 tenrecs (table 1). We recognised that our landmarks and curves for the  
156 mandibles focus particular attention on the ascending ramus (condyloid,  
157 condylar and angular processes, figure ??). Therefore we deleted the three  
158 semilandmark curves around these structures and repeated our disparity  
159 calculations. In this case we found no significant differences in disparity  
160 between the two Families (table 1). Therefore, our results seem to indicate  
161 that golden moles have greater morphological variation in the posterior  
162 structures of their mandibles compared to tenrecs.

163 Given that these posterior structures act as muscle attachment and



164 articulation sites for connections with the upper jaw, one might expect  
165 that golden moles with highly disparate posterior mandible morphologies  
166 should also show high variability in the corresponding mandible  
167 articulation areas of the skull. However, we could not locate reliable,  
168 homologous points accurately on those areas of the skull pictures in  
169 lateral view. Instead, our landmarks and semilandmark curves for the  
170 skulls in lateral view focus attention on morphological variation in the  
171 dentition and the overall shape of the top and back of the skulls (figure  
172 ??). This may explain why golden mole skulls in lateral view do not show  
173 the same pattern of higher disparity compared to tenrecs that we see in  
174 our analyses of the mandibles. However, further investigation is required  
175 to identify possible reasons why golden moles appear to show such  
176 variation in the posterior structures of their mandibles.

177 We used variation in skull and mandible shapes as proxy measures for  
178 overall morphological diversity within the two Families. Many other  
179 studies also use skulls to study phenotypic variation within species  
180 (Blagojević & Milošević-Zlatanović, 2011; Bornholdt et al., 2008), to  
181 delineate species boundaries within a clade (e.g. Panchetti et al., 2008) or  
182 for cross-taxonomic comparative studies of phenotypic (dis)similarities  
183 (e.g. Ruta et al., 2013; Goswami et al., 2011; Wroe & Milne, 2007).

184 However, studies of morphological disparity are inevitably constrained  
185 to measure diversity within specific traits rather than overall phenotypes  
186 (Roy & Foote, 1997). Disparity calculations based on skull shape can yield  
187 similar results compared to analyses of whole-skeleton discrete characters  
188 and limb proportion data sets (Foth et al., 2012). Yet it is still possible that  
189 comparing disparity in tenrecs and golden moles using non-cranial  
190 morphological measures could produce different results. For example,

tenrecs inhabit a wide variety of ecological niches and habitats including terrestrial, arboreal, semi-aquatic and semi-fossorial environments (Soarimalala & Goodman, 2011). In contrast, although golden moles occupy a wide altitudinal, climatic and vegetational spectrum of habitats (Bronner, 1995), they are all fossorial species which, superficially at least, appear to be less functionally diverse than tenrecs. Therefore, comparing the disparity of limb morphologies within the two Families could indicate that tenrecs are more morphologically diverse than golden moles and therefore support the claim that tenrecs are an exceptionally diverse group.

Our analyses are the first measures of morphological diversity within tenrecs, a group which is commonly cited as an example of an adaptive radiation (Olson, 2013). Evidence of exceptional morphological diversity is one criterion for designating a clade as an adaptive radiation (Losos & Mahler, 2010). However, we found that tenrecs are no more morphologically diverse than their closest relatives and therefore, within our tests, do not appear to show the exceptional diversity which characterises an adaptively radiated group.

The evolution of cranial shape (both upper skull and mandible), particularly dental morphology, has obvious correlations with dietary specialisations and occupation of specific ecological niches (e.g. Wroe & Milne, 2007). Considering the wide ecological diversity of the tenrec Family; semi-fossorial, arboreal, terrestrial and semi-aquatic (Soarimalala & Goodman, 2011), we think that it is reasonable to expect that this variety should be reflected in skull morphology. However, we have not included any measures of the 'adaptiveness' of cranial shape in our analyses and therefore our analyses should not be considered to be an

218 explicit test of whether or not tenrecs are an adaptive radiation (Losos &  
219 Mahler, 2010). Instead we have made the first step towards understanding  
220 the apparent phenotypic diversity within tenrecs within a quantitative  
221 framework. Future work should focus on explicit measures of the  
222 'adaptiveness' and functional importance of tenrec cranial and  
223 post-cranial morphologies to understand the significance of  
224 morphological diversity within the Family (e.g. Mahler et al., 2010).  
225 However, we also recognise that strict, statistically based categorisations of  
226 clades as being adaptive radiations or not are not always biologically  
227 meaningful or helpful when it comes to trying to understand patterns of  
228 phenotypic diversity (Olson & Arroyo-Santos, 2009).

229 We have presented the first quantitative study which tests the common  
230 claim that tenrecs are an exceptionally diverse group (Olson, 2013;  
231 Soarimalala & Goodman, 2011; Eisenberg & Gould, 1969). Focusing on  
232 cranial diversity is only one aspect of morphological variation and further  
233 analyses are required to test whether other morphological traits yield  
234 similar patterns. However, our results provide a clear indication that  
235 phenotypic variety within tenrecs is perhaps not as exceptional as it first  
236 seems.

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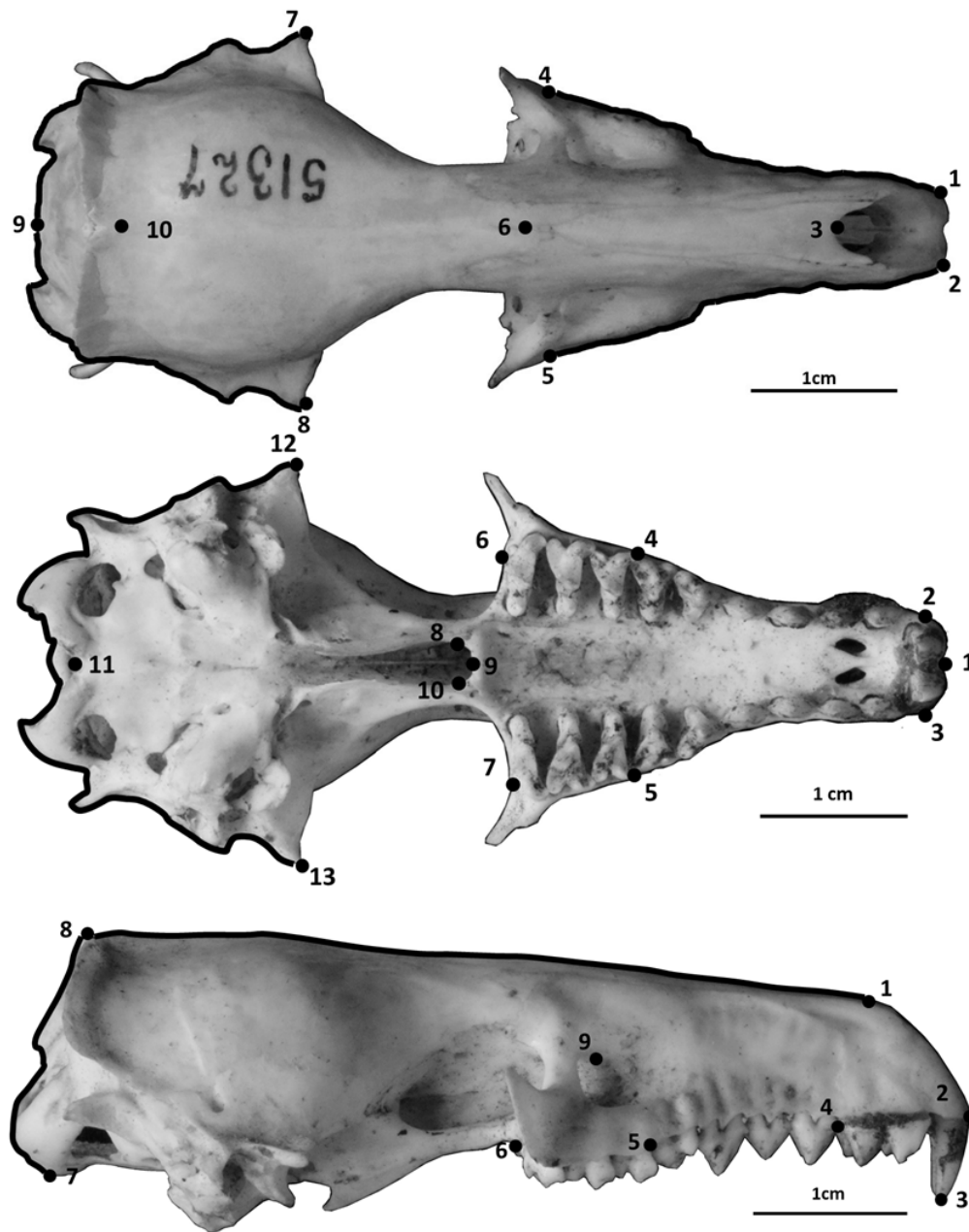


Figure 1: Landmarks (numbered points) and curves (black lines) used to capture the morphological shape of skulls in dorsal, ventral and lateral views respectively. Curves were re-sampled to the same number of evenly-spaced points. See Supplementary Material for descriptions of the curves and landmarks. The specimens belong to two different *Potamogale velox* (Tenrecidae) skulls: accession number AMNH 51327 (dorsal) and BMNH 1934.6.16.2 (ventral and lateral)

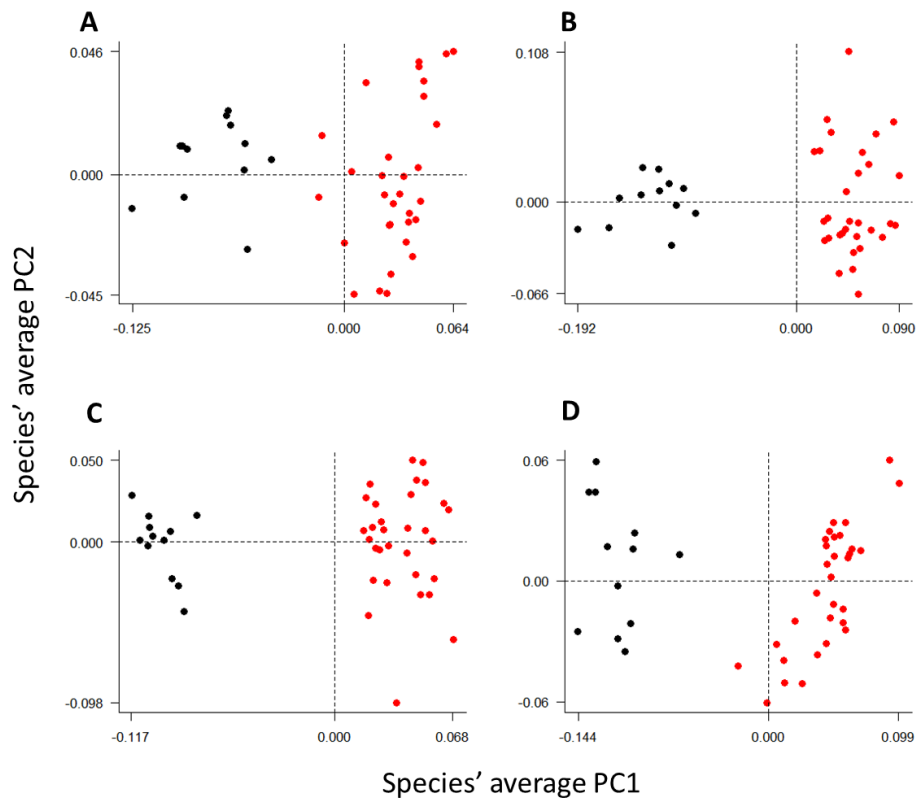


Figure 2: Principal components plots of the morphospaces occupied by tenrecs (red,  $n = 31$  species) and golden moles (black,  $n = 12$ ) for the skulls: dorsal (A), ventral (B), lateral (C) and mandibles (D) analyses. Axes are PC1 and PC2 of the average scores from a PCA analysis of mean Procrustes shape coordinates for each species.

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Table 1: Disparity comparisons between tenrecs (T) and golden moles (G) for each of our data sets(rows) and four disparity metrics (columns). ‘Mandibles:one curve’ refers to our shape analysis of mandibles excluding the three curves around the posterior structures of the jaw (figure ??). Significant differences are highlighted in bold with the corresponding p value in brackets. Disparity metrics are: sum of variance, product of variance, sum of ranges and product of ranges

<b>Disparity metric</b>	<b>SumVar</b>	<b>ProdVar</b>	<b>SumRange</b>	<b>ProdRange</b>
Skulls dorsal	T>G	T>G	T>G	T>G
Skulls lateral	T>G	T>G	T>G	T>G
Skulls ventral	T>G	G>T	T>G	T>G
Mandibles	G>T	<b>G&gt;T* (0.008)</b>	<b>T&gt;G* (0.025)</b>	<b>G&gt;T* (0.009)</b>
Mandibles:one curve	G>T	G>T	T>G	T>G

Table 2: Disparity comparisons between non-*Microgale* tenrecs (T) and golden moles (G) for each of our data sets(rows) and four disparity metrics (columns). Significant differences are highlighted in bold with the corresponding p value in brackets. Disparity metrics are; sum of variance, product of variance, sum of ranges and product of ranges.

<b>Disparity metric</b>	<b>SumVar</b>	<b>ProdVar</b>	<b>SumRange</b>	<b>ProdRange</b>
Skulls dorsal	T>G	T>G	T>G	T>G
Skulls lateral	<b>T&gt;G* (0.014)</b>	T>G	<b>T&gt;G* (0.001)</b>	<b>T&gt;G*(0.003)</b>
Skulls ventral	T>G	T>G	T>G	T>G
Mandibles	T>G	G>T	T>G	G>T

Table 3: npMANOVA comparisons of morphospace occupation for tenrecs and golden moles in each of the four analyses (three views of skulls and mandibles). In each case the two families occupy significantly different areas of morphospace.

<b>Analysis</b>	<b>F</b>	<b>R<sup>2</sup></b>	<b>p value</b>
Skulls dorsal	66.02	0.62	0.001
Skulls ventral	100.74	0.71	0.001
Skulls lateral	75.07	0.65	0.001
Mandibles	59.34	0.59	0.001