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ORIGINAL ARTICLE

Distinctive features of the skull of the Ryukyu Scops Owl from Minami-daito Island, revealed by computed tomography scanning

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Abstract Morphological differentiation of island-dwelling organisms provides model systems for studying evolution. Computed tomography (CT) scanning is an entirely non-destructive technique that provides detailed three-dimensional (3D) images of physical structures. Geometric morphometrics has been increasingly used in avian morphology studies by analyzing 3D data obtained from CT scans. We used geometric morphometrics to evaluate the morphological details of the skulls of three, genetically distinct, island populations of the Ryukyu Scops Owl *Otus elegans*: *O. e. elegans* from the northern part of the Ryukyu Archipelago, *O. e. elegans* from the southern part of the Ryukyu Archipelago, and *O. e. interpositus* from Minami-daito Island. Skulls were scanned using an X-ray CT system and the digitized 3D coordinates of 16 landmarks for each skull were analyzed in order to describe geometric morphometric features. *O. e. interpositus* was found to have a significantly smaller skull than either population of *O. e. elegans*. From principle component analysis of shape variation, we also found that the skull shape of *O. e. interpositus* differed significantly from both the northern and southern groups of *O. e. elegans*. This difference was in terms of PC1, which mainly represented relative anteroposterior length, and angle of the orbit. We inferred that the small skull of *O. e. interpositus* is partly a consequence of the particular founders of the population, or evolutionary selection that has taken place on Minami-daito Island and that the distinctive shape of the skull of *O. e. interpositus* is partly a consequence of adaptations for foraging efficiency, or of morphological integration.

Key words CT scan, Geometric morphometrics, island, *Otus elegans*, skull

Morphological features of organisms can reflect their evolutionary histories. In addition, population differentiation between islands can provide model systems for studying evolution (Santos et al. 2016). Classic studies of evolutionary biology have focused on the morphology of island birds, such as Darwin's finches (Geospizinae) (Grant & Grant 2011) and the Hawaiian honeycreepers (Drepanididae) (Pratt 2005). Herrel et al. (2007) suggested that the internal cranial skeleton has crucial importance in both ecological and behavioral contexts. Therefore, we considered

that evaluating the differentiation of skulls between populations of island birds has the potential for providing insights into the ways in which ecological and behavioral factors affect evolution.

The Ryukyu Scops Owl *Otus elegans* is an island species with a very restricted range of no more than 1,000 km from north to south. It is mainly distributed throughout the Ryukyu Archipelago of Japan, a small island off southeast Taiwan, and south to small islands north of Luzon, Philippines (Ornithological Society of Japan 2012; Takagi et al. 2015; Gill & Donsker 2017). Four subspecies are currently recognised: *O. e. elegans*, *O. e. interpositus*, *O. e. botelensis*, and *O. e. calayensis* (Gill & Donsker 2017). Two

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subspecies occur in Japan: nominate *O. e. elegans*, which occurs widely on the continental islands of the Ryukyu Archipelago, and the isolated *O. e. interpositus*, which is endemic to oceanic Minami-daito Island (Fig. 1).

The current subspecific classification is mainly based on plumage characteristics (Kuroda 1923), but more recently, Takagi (2011) revealed that the three populations, those on island groups north and south of the Kerama Gap, and those on Minami-daito Island (Fig. 1) differed in their vocalisations. Features of their morphology seem to be largely attributable to their geohistorical isolation. A recent mitochondrial cytochrome c oxidase subunit I (COI) analysis has supported the subdivision of *O. e. elegans* into northern and southern populations (Saitoh et al. 2015). The genetic independence of the Minami-daito Island *O. e. interpositus* population has also been revealed by a STRUCTURE analysis (Pritchard et al. 2000) of microsatellite regions (Takagi & Saitoh in preparation). Although their plumage, acoustic, and genetic characteristics have been described in the previous studies mentioned above, their internal morphology has never been described.

We used landmark-based geometric morphometrics to evaluate morphological variation in Computed tomography (CT) scanned skulls of the Ryukyu Scops Owl. CT scanning has become increasingly popular among natural science disciplines as it is entirely non-destructive and provides detailed three-dimensional (3D) images of structures (Matthews & Plessis 2016). The CT images allow us to manipulate, section, dissect and measure internal morphology as well as external morphology (Abel et al. 2012). This characteristic of CT scanning is a necessary property for obtaining internal morphological data of objects that must not be damaged or destroyed, such as precious museum specimens.

Landmark-based geometric morphometrics has been increasingly applied in a range of ornithological studies, such as those on the relationships between ecology and morphology (Kulemeyer et al. 2009; Bright et al. 2016; Matsui et al. 2016) and avian evolution (Bhullar et al. 2012; Bhullar et al. 2015). As traditional morphometrics is based on multivariate analysis of focused variables, such as linear measurements, ratios, angles, areas, and volumes, the results are restricted to those relevant to these

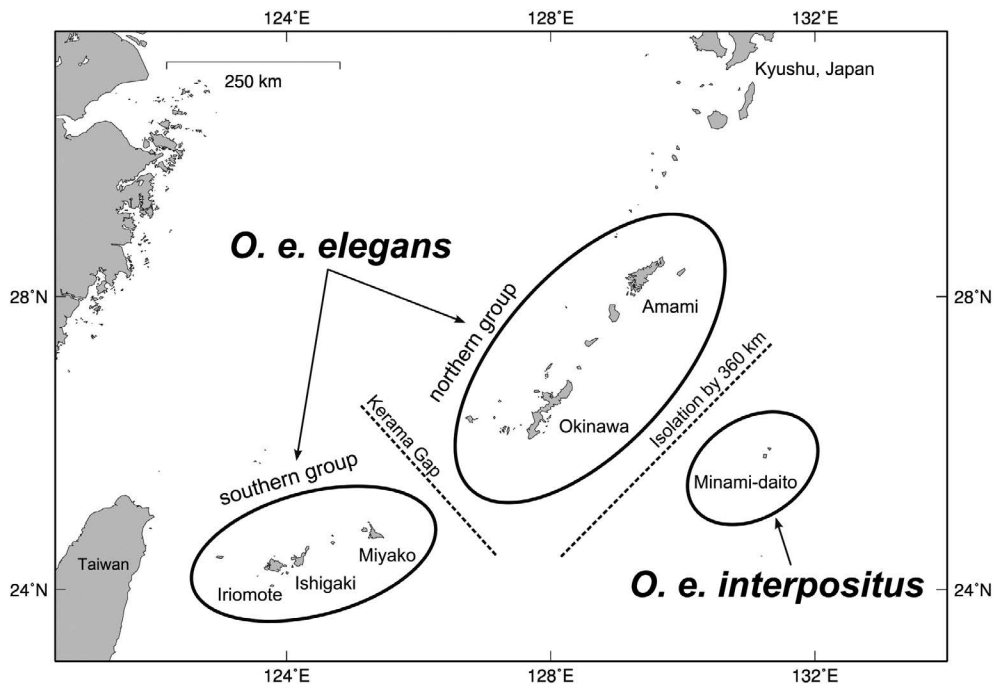


Fig. 1. Three groups of the Ryukyu Scops Owl, the northern and southern groups, and Minami-daito Island, based on genetic analyses (Saitoh et al. 2015; Takagi & Saitoh in preparation). Ovals show the three groups; the dotted line indicates the Kerama Gap and isolation of Minami-daito Island. The Kerama Gap geographically divides subspecies *O. e. elegans* into northern and southern groups. *O. e. interpositus*, isolated on Minami-daito Island, is separated by 360 km from Okinawa, the nearest island.

a priori selected variables and their combinations (Mitteroecker & Gunz 2009). However, as landmark-based geometric morphometrics analyze Cartesian landmark coordinates (Mitteroecker & Gunz 2009), results can be drawn from any possible relative positioning of the landmarks. Therefore, landmark-based geometric morphometrics is superior to traditional morphometrics, in terms of finding unexpected morphological differences.

In this study, we explored the skull morphology of three different populations of the Ryukyu Scops Owl in order to establish whether any characteristics differ among them, and sought explanations for the differences we found.

MATERIALS AND METHODS

A total of 28 specimens were derived as follows: six (two males, one female, three of unknown sex) from the Ryukyu Archipelago north of the Kerama Gap; seven (two males, five of unknown sex) from the Ryukyu Archipelago south of the Kerama Gap; and 15 (seven males, four females, four of unknown

sex) from Minami-daito Island. Twenty of the specimens were dry museum skins and eight were frozen bodies; 20 were adult, one was a yearling, and seven were of unknown age. All of the specimens of unknown age were sufficiently ossified to allow CT imaging: they were considered to be either yearlings or adults, not juveniles (see Appendix 1).

All 28 skulls were scanned using an X-ray CT system (La Theta LTC-100, Hitachi Aloka Medical, Tokyo, Japan) with slice thickness set to 0.2 mm. Superficial models of the skulls were reconstructed from their CT-data. The digitized 3D-coordinates of the 16 landmarks on the surfaces of model skulls were acquired by means of ImageJ 1.49 (Schneider et al. 2012) (Fig. 2, Table 1). As avian adult neurocranium bones are fused (Zusi 1993), it proved difficult to place landmarks on skulls. Furthermore, some specimens had been damaged around the *foramen magnum* by taxidermists removing brain tissue from the cranium, making it difficult for us to put landmarks on the skull images. Accordingly, we chose 16 landmarks as a compromise between maximizing the number of landmarks and avoiding missing data,

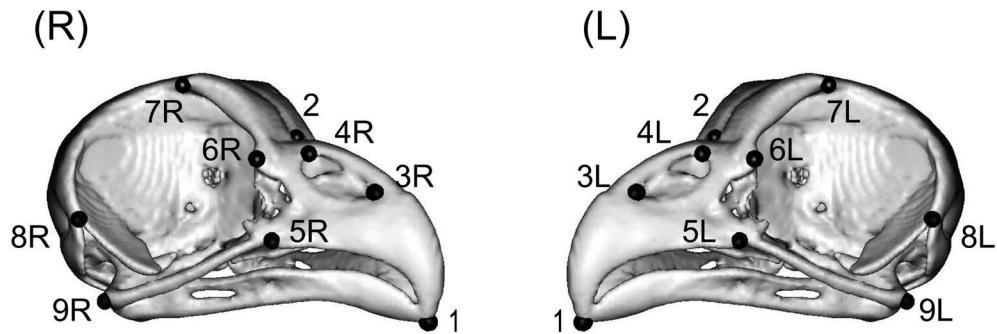


Fig. 2. Landmarks (black dots) on right (R) and left (L) hemispheres of a Ryukyu Scops Owl skull.

Table 1. Landmarks on the skull of the Ryukyu Scops Owl.

Landmark No.	description
1	tip of premaxillary
2	midpoint of craniofacial hinge
3L/3R	maximum curvature point of anterior end of left/right nares
4L/4R	maximum curvature point of posterior end of left/right nares
5L/5R	left/right attachment point of jugal and maxillary
6L/6R	outermost point of left/right lacrimal
7L/7R	tip of left/right dorsal process on orbit
8L/8R	diverging point of left/right postorbital process and tympanic wing
9L/9R	outermost point of left/right quadrate

Description follows nomenclature used in Zusi (1993). Landmark No. corresponds to Fig. 2.

which emerged in the data of damaged specimens when we put landmarks around the *foramen magnum*. The geometric features of the skulls were summarized into landmark configurations.

In order to describe the geometric morphometric features of the skulls, first we captured size and shape variation from landmark configurations. Second, we compared skull size and shape between the three groups. Finally, we examined whether the differences found were explained by allometry or not.

Size and shape information were extracted from the original landmark dataset using Generalized Procrustes Analysis (GPA). GPA is based on a least-square approach that minimizes the distance between homologous landmarks by translating, scaling, and rotating the original landmark configurations (Slice 2007). The scaling factor extracted through GPA is called “centroid size”. The coordinates of the resulting centered, scaled, and rotated landmark configurations are called “Procrustes shape coordinates”. Size was represented in centroid size and shape by Procrustes shape coordinates. GPA was conducted using the *ProcGPA* function, implemented in R version 3.1.3 (R Core Team 2015) with the package “Morpho” (Schlager 2015).

To evaluate the significance of differences in skull centroid size, we used the Kruskal-Wallis test and the pairwise Wilcoxon rank sum test. Principal component analysis (PCA) summarized variation in Procrustes shape coordinates. First, major principal components (PC), which explained more than 80% of total variance, were compared by MANOVA to compare PC scores among three groups. Subsequently, the significance of the difference in each PC score was assessed by ANOVA and pairwise t test if MANOVA indicated significance. Holm’s correction for P-value adjustment (Holm 1979) was used for all ANOVAs and pairwise comparisons. Before all intergroup comparisons were made, we assessed sexual dimorphism using specimens from Minami-daito Island, as there were multiple samples of both sexes. As no sexual dimorphism either in centroid size or first major PCs of Procrustes shape coordinates was detected (see results), we conducted all analyses on pooled data without separating for sex. We also conducted all analyses without separating for age because all samples including unknown age were deemed to be yearling or adult. Sufficient ossification of yearlings, combined with knowing that the owls first commence breeding when yearlings, support the fact that yearlings have a full grown skeleton and

do not differ from adults. We excluded samples of juveniles because they may have different skull structure (Hogg 1980). Visualization of the deformation of skulls along PC axes was obtained from thin plate spline interpolation (Mitteroecker & Gunz 2009). All statistical analyses were conducted with R version 3.1.3. Calculation of thin plate spline interpolation was done using the *tps3d* function implemented in the R package “Morpho” (Schlager 2015). Deformed skulls were visualized and captured using MeshLab version 1.3.3 (Cignoni et al. 2008).

Explaining shape variation between the three groups by allometry requires one regression line that relates shape to size for the merged group. For the purpose of drawing this regression line, we adopted a procedure consisting of two steps. The first step involved fitting a regression line for each of the three groups. Depending on the results of the first step, we followed one of the following three sub-procedures in the second step: a) If no significant regression line was detected in any of the three groups, or detected for only one group, we conducted a regression analysis using merged data; b) If significant regression lines were obtained for at least two of the groups and these lines did not differ statistically from each other, we also conducted merged data analysis; c) If regression lines for at least two of the groups proved to be significantly different, pooled analysis was not justifiable, and not conducted.

We conducted Procrustes ANOVA (Goodall 1991) using the *procD.lm* function of R package “geomorph” (Adams & Otárola-Castillo 2013) in order to test the significance of regressions with a permutation test in the first step. In cases when significant regression lines were detected (i.e., cases (b) or (c) above), we aimed to test the homogeneity of slopes and intercepts using the *advanced.procD.lm* function of R package “geomorph”. To summarize variation in the Procrustes shape coordinates, which was not explained by the size factor, we performed PCA on the residual of the regression analysis of merged data. We used the same statistical procedures for analysing residual Procrustes shape coordinates as described above.

RESULTS

Size difference

Centroid size did not differ significantly between males and females (Wilcoxon rank sum test, $n_1=7$, $n_2=4$, $W=19$, NS), but differed significantly between

the three population groups (Kruskal-Wallis test, chi-squared=10.58, df=2, $P<0.01$). Pairwise Wilcoxon rank sum tests revealed that the centroid size of *O. e. interpositus* was significantly smaller than

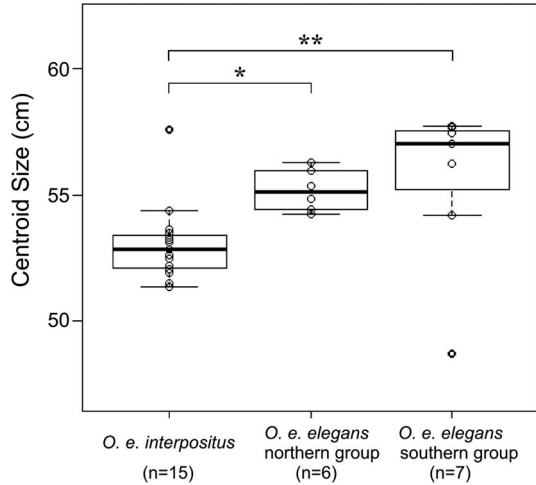


Fig. 3. Comparison of centroid size between three groups of Ryukyu Scops Owls. * $P<0.05$, ** $P<0.01$

that of the northern ($n_1=15$, $n_2=6$, $W=83$, adjusted $P<0.05$, Fig. 3) and that of the southern ($n_1=15$, $n_2=7$, $W=85$, adjusted $P<0.05$, Fig. 3) groups of *O. e. elegans*. However, differences between the two groups of *O. e. elegans* ($n_1=6$, $n_2=7$, $W=29$, NS, Fig. 3) were not significant.

Shape difference

The first eight PCs in the original Procrustes shape coordinates explained 81.3% of the total shape variation (see Appendix 2 for detailed data). These PC scores did not differ significantly between males and females (MANOVA, df=8, 2, $F=2.85$, Pillai Trace=0.92, NS), but did differ significantly between the three regional groups (MANOVA, df=16, 38, $F=2.96$, Pillai Trace=1.11, $P<0.01$). PC1, which explained 22.9% of total shape variation, showed a significant difference among the three groups (ANOVA, df=2, 25, $F=39.32$, $P<0.01$, Appendix 2, Fig. 4). With the exception of PC1, none of the other major PCs differed significantly between the

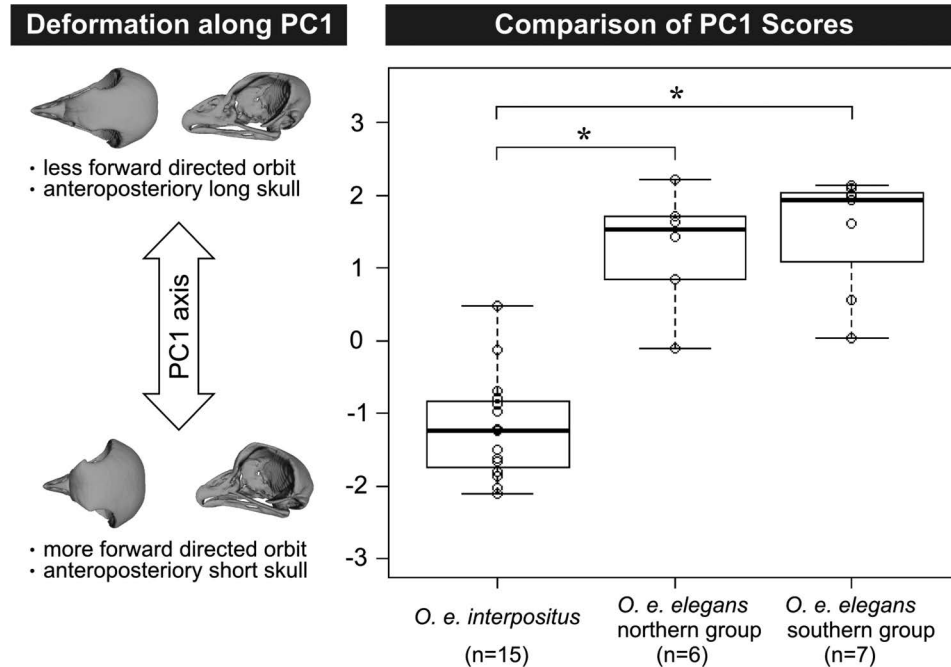


Fig. 4. Comparison of PC1 score of original Procrustes shape coordinates and deformation of the skull along the PC1 axis in the Ryukyu Scops Owl. * $P<0.001$. Landmarks on average shaped skulls were moved along the PC1 axis so that the resultant landmark configuration has a mean +4SD PC1 score (upper row) or mean -4SD PC1 score (lower row). Each skull is expressed as a vector whose components are coordinates of landmarks on the skull. PC score consists of a linear combination of coordinates of landmarks. Therefore, moving landmarks on a skull, i.e. adding a vector to the vector of coordinates of landmarks, causes changes in PC scores of the skull. Actually, adding a scaled eigenvector of PCA can increase or decrease the corresponding PC score. Extent of the change depends on the scaling factor (see Appendix of Mitteroecker et al. 2013). The remaining surface of an average shaped skull was deformed by thin plate spline interpolation in accordance with the changes in the landmark positions. Deformed skulls are shown from lateral (right) and dorsal views (left).

three groups (see Appendix 2 for detailed results of ANOVA). Further analysis was conducted only for PC1. The PC1 score of *O. e. interpositus* was significantly smaller than those of the northern (t-test, $df=8.36$, $t=6.51$, $P<0.001$) and the southern groups (t-test, $df=10.37$, $t=7.28$, $P<0.001$). The northern and southern groups of *O. e. elegans* did not differ from each other (t-test, $df=10.79$, $t=0.43$, NS). Deformation of the average-shaped skull along the PC1 axis showed that PC1 mainly represented relative anteroposterior length and the angle of the orbit (Fig. 4). In other words, birds belonging to *O. e. interpositus* have relatively short skulls with orbits positioned further forward than in other populations.

Allometry

We did not detect any significant regression lines for the three groups by multivariate regression analysis of Procrustes shape coordinates on log centroid size (permutation test; 10,000 permutations, NS for all groups). We obtained a significant regression line when analyzing the merged data (permutation test; 10,000 permutations, $P<0.001$). The first eight PCs from the PCA based on residual Procrustes shape coordinates explained 80.5% of the total shape variation (see Appendix 3 for detailed data). As none of these PC scores differed significantly between the three groups (MANOVA, $df=16$, 38 , $F=1.80$, Pillai Trace=0.86, NS), we did not perform any subsequent ANOVAs for each PC.

DISCUSSION

The objective of this research was to study, detect and evaluate any differences between the skulls of three geographically separated populations of the Ryukyu Scops Owl, to infer causal factors for any differences, and to propose explanations for how differences might have evolved. Analyses revealed differences both in size and shape between *O. e. interpositus* and *O. e. elegans*. Multivariate regression of Procrustes shape coordinates on the logarithm of centroid size for each group did not indicate significant allometric relationships. However, regression analysis of a merged group provided significance. Analyses based on residual Procrustes shape coordinates did not indicate subspecific shape differences.

Pairwise comparisons revealed that centroid size differed between *O. e. interpositus* and *O. e. elegans*. We inferred that small skull size on the small (ca. 30 km²), isolated, oceanic Minami-daito Island might

be partly attributable to founder effects. A STRUCTURE analysis of microsatellite regions (Takagi & Saitoh in preparation) suggests that the population on Minami-daito Island was founded by individuals from the population on Okinawa Island, which is included within the northern group of *O. e. elegans*. Our results showed that some individual *O. e. elegans* had skulls as small as those of *O. e. interpositus* (Fig. 3), and we propose that founders reaching Minami-daito Island may have had such characteristics. Conversely, however, other evolutionary selection processes may have favored small skull size on Minami-daito Island after initial arrival. Clegg and Owens (2002) discussed how the effects of changes in feeding ecology, intraspecific competition, energetic constraints and physiological optimization in birds on small islets seem to have an important role in size evolution. However, as yet we do not have data supporting or confirming either perspective.

Principle component analysis revealed that cranial shape also differed substantially between *O. e. interpositus* and *O. e. elegans*, for which there seem to be three possible explanations. The first possibility is allometry, one of the factors causing morphological variation among organisms that differ in body size (Mitteroecker et al. 2013). The results of the PCA of residual Procrustes shape coordinates suggest the possibility that variation between subspecies can be simply explained as an allometric effect of skull size. The disappearance of significance of PC1, after controlling for size, suggest that the PC1 of original Procrustes shape coordinates mainly represents allometric variation in the data. Actually, allometry is the most dominant factor of shape and form variation (Mitteroecker et al. 2013). However, explanation by allometry requires careful interpretation of results, because multivariate regressions in each group did not indicate significant allometric relationships. If each group actually has a different regression line, then multivariate regression analysis for merged data is not justified and we need a different explanation, such as that offered below.

The second possibility is that adaptation to foraging behaviour using binocular vision has occurred. Orientation of the orbits may be related to variation in the visual field (Heesy 2004; Fernández-Juricic et al. 2008), which may reveal a trade-off relationship between foraging and predator detection (Martin 2014). Hunting owls swing their feet up into the area of binocular vision just prior to prey capture (Martin 2007). Although forward positioned eyes

increase accuracy during this prey capturing behavior, such positioning decreases lateral vision rendering the hunting owl more susceptible to predators attacking from the side. Potential nocturnal predators, such as *Protobothrops* snakes, do not occur on oceanic Minami-daito Island, whereas they do occur on almost all of the other continental islands in the Ryukyu Archipelago. Adaptations among the owls on Minami-daito Island might have led to them having eyes positioned further forward thus increasing their hunting accuracy. To verify this trade-off relationship, future studies should include analysis of morphometric data in relation to the presence or absence of predators on each island. Quantitative evaluation of visual fields is also desirable (i.e., Fernández-Juricic et al. 2008; Martin & Piersma 2009).

The third explanation for the difference in skull shape is morphological integration. Each part of the skull is integrated with other parts because they develop, function, and evolve together (Klingenberg 2013). The observed shape difference may result from such covariation of skull parts. In other words, part of the observed variation may be caused by variation in other parts, i.e. variation that is explained by other factors such as adaptation. In this study, we were compelled to adopt a small number of landmarks due to the limitation of the material properties of the specimens (as described above). To accurately evaluate shape covariation among parts, many more landmarks including semi-landmarks are needed (Kulemeyer et al. 2009; Gunz & Mitteroecker 2013).

Our study has shown that two subspecies of Ryukyu Scops Owl exhibit substantially different skull size and shape. Such differences may be explained by allometry, adaptation, or morphological integration. However, the variation detected should be interpreted carefully. Our results are based on a relatively small sample size including individuals of unknown sex. A larger sample with sexed specimens would help to improve our understanding of their morphological evolution.

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Skull features in the Ryukyu Scops Owl

Appendix 1. Ryukyu Scops Owl specimens examined in this study.

Specimen No.	Owner	Island	Subspecies	Group	Sex	Stage	State
YIO-19205	YIO	Minami-daito	<i>interpositus</i>		Female	Unknown	Museum skin
YIO-19207	YIO	Minami-daito	<i>interpositus</i>		Male	Unknown	Museum skin
YIO-19208	YIO	Minami-daito	<i>interpositus</i>		Female	Unknown	Museum skin
YIO-19210	YIO	Minami-daito	<i>interpositus</i>		Male	Unknown	Museum skin
YIO-19216	YIO	Okinawa	<i>elegans</i>	northern	Male	Adult	Museum skin
YIO-19229	YIO	Iriomote	<i>elegans</i>	southern	Unknown	Adult	Museum skin
YIO-62371	YIO	Ishigaki	<i>elegans</i>	southern	Unknown	Unknown	Museum skin
YIO-64984	YIO	Iriomote	<i>elegans</i>	southern	Unknown	Adult	Museum skin
2004-0129	YIO	Iriomote	<i>elegans</i>	southern	Unknown	Unknown	Frozen body
2015-1084	YIO	Minami-daito	<i>interpositus</i>		Female	Adult	Museum skin
2015-1086	YIO	Minami-daito	<i>interpositus</i>		Male	Adult	Museum skin
2015-1087	YIO	Minami-daito	<i>interpositus</i>		Unknown	Adult	Frozen body
2015-1089	YIO	Minami-daito	<i>interpositus</i>		Unknown	Adult	Frozen body
2015-1092	YIO	Minami-daito	<i>interpositus</i>		Unknown	Adult	Frozen body
2015-1093	YIO	Minami-daito	<i>interpositus</i>		Female	Yearling	Museum skin
2015-1094	YIO	Minami-daito	<i>interpositus</i>		Male	Adult	Frozen body
2015-1104	YIO	Minami-daito	<i>interpositus</i>		Unknown	Adult	Frozen body
2015-1105	YIO	Minami-daito	<i>interpositus</i>		Male	Adult	Museum skin
2015-1106	YIO	Minami-daito	<i>interpositus</i>		Male	Adult	Frozen body
2015-1110	YIO	Minami-daito	<i>interpositus</i>		Male	Adult	Frozen body
NSMTA-13193	NMNS	Iriomote	<i>elegans</i>	southern	Unknown	Unknown	Museum skin
NSMTA-17395	NMNS	Okinawa	<i>elegans</i>	northern	Male	Adult	Museum skin
NSMTA-17402	NMNS	Okinawa	<i>elegans</i>	northern	Female	Adult	Museum skin
NSMTA-17501	NMNS	Amami	<i>elegans</i>	northern	Unknown	Adult	Museum skin
NSMTA-17504	NMNS	Amami	<i>elegans</i>	northern	Unknown	Adult	Museum skin
NSMTA-17511	NMNS	Amami	<i>elegans</i>	northern	Unknown	Adult	Museum skin
NSMTA-17615	NMNS	Iriomote	<i>elegans</i>	southern	Male	Adult	Museum skin
NSMTA-17616	NMNS	Iriomote	<i>elegans</i>	southern	Male	Adult	Museum skin

Abbreviations: YIO, Yamashina Institute for Ornithology; NMNS, National Museum of Nature and Science.

Appendix 2. Results of the principal component analysis (PCA) on original Procrustes shape coordinates and the analysis of variance (ANOVA).

Principle component	Explained variance (%)	Cumulative explained variance (%)	ANOVA		
			F _(2, 25)	P	Significance
PC1	22.9	22.9	39.32	0.000	*
PC2	16.6	39.5	0.01	0.995	
PC3	13.8	53.3	0.46	0.637	
PC4	8	61.3	3.31	0.053	
PC5	7	68.3	0.51	0.608	
PC6	5.4	73.7	0.06	0.947	
PC7	4.3	78.1	0.18	0.838	
PC8	3.2	81.3	0.64	0.536	

Significance: * P<0.01 after Holm's P-value correction

Appendix 3. Results of the principal component analysis (PCA) on residual Procrustes shape coordinates and the analysis of variance (ANOVA).

Principle component	Explained variance (%)	Cumulative explained variance (%)	ANOVA		
			F (2, 25)	P	Significance
PC1	19.8	19.8	2.98	0.069	
PC2	15.8	35.6	2.43	0.108	
PC3	14.1	49.7	2.53	0.100	
PC4	8.9	58.6	3.44	0.048	
PC5	7.9	66.5	0.51	0.606	
PC6	5.6	72.1	0.22	0.807	
PC7	4.8	76.9	0.23	0.797	
PC8	3.6	80.5	0.68	0.519	

Significance: * $P < 0.01$ after Holm's P-value correction