RESEARCH ARTICLE

In and Out the Amazonia: Evolutionary Ecomorphology in Howler and Capuchin Monkeys

Carlo Meloro • Nilton Cáceres • Francesco Carotenuto • Jonas Sponchiado • Geruza Leal Melo • Federico Passaro • Pasquale Raia

Received: 28 March 2013 / Accepted: 26 June 2013 / Published online: 12 July 2013 - Springer Science+Business Media New York 2013

Abstract The impact of environmental variation on phenotypic diversification is one major issue in evolutionary studies. Environmental variation is thought to be a primary factor in evolution, especially at high latitudes. In contrast, tropical areas are traditionally viewed as the cradle where the long-term effects of biological interactions on phenotypic change reside. We analyse patterns of skull shape variation in two New World monkey groups: capuchins and howlers. These two monophyletic clades are exceptionally similar in terms of the geographic distribution of their species. Yet, their body size and diet are different: howler monkeys are large and almost exclusively folivorous, whereas capuchins are small omnivorous. We found that the size, and direction of vectors of phenotypic changes across South American biomes in those clades are not statistically

Electronic supplementary material The online version of this article (doi:[10.1007/s11692-013-9244-5\)](http://dx.doi.org/10.1007/s11692-013-9244-5) contains supplementary material, which is available to authorized users.

C. Meloro (⊠) · F. Carotenuto · F. Passaro · P. Raia Dipartimento di Scienze della Terra, Universita` degli Studi di Napoli 'Federico II', L.go San Marcellino 10, 80138 Naples, Italy e-mail: carlo.meloro@hyms.ac.uk

F. Carotenuto e-mail: f.carotenuto@ymail.com

F. Passaro e-mail: federico.passaro@unina.it

P. Raia e-mail: pasquale.raia@libero.it

N. Cáceres

Department of Biology, Laboratory of Ecology and Biogeography, CCNE, Federal University of Santa Maria, Santa Maria, RS 97110-970, Brazil e-mail: niltoncaceres@gmail.com

different. This similarity persists after removing the strong impact of allometry in our data. Additionally, partial least squares and comparative analyses confirm that ''allometry free'' skull shape is influenced to the same set of environmental variables in both clades. This study remarks the paramount importance of both body size and environmental variation on phenotypic evolution.

Keywords Skull shape - Climatic adaptation - Partial least squares · South America · Amazon forest

Introduction

When climatic change occurs, species respond by adapting, shifting their geographic distribution to track their preferred habitats, or going extinct (Davis et al. [2005;](#page-12-0) Raia et al. [2012\)](#page-13-0). Adaptation as driven by environmental variation is one of the common mechanisms underlying

J. Sponchiado

G. L. Melo

Programa de Pós-Graduação em Ecologia e Conservação, CCBS, Universidade Federal do Mato Grosso do Sul, Cx.P. 549, Campo Grande, MS 79070-900, Brazil e-mail: geruzalm@yahoo.com.br

Department of Biology, Programa de Pós-Graduaçao em Biodiversidade Animal, CCNE, Federal University of Santa Maria, Santa Maria, RS 97110-970, Brazil e-mail: jsponchiado@yahoo.com.br

phenotypic evolution. The formation of geographic barriers, expansion and contraction of biomes in relation to climatic change, and secular trends in global temperature are some of the many ''environmental'' factors known to prompt phenotypic diversification (Rundell and Price [2009](#page-13-0); Hoffmann and Sgrò [2011;](#page-12-0) Raia et al. [2013\)](#page-13-0). Within this context, tropical areas always received special attention for the high diversity of species and ecomorphotypes they harbor. As the intensity of environmental variation is directly linked to latitude (Davies et al. [2009\)](#page-12-0) species in the tropics are expected to be less sensitive to climatic change. In fact, the high diversity of species and ecomorphs in the tropics is traditionally attributed to high ecosystem resilience and primary productivity (Currie [1991](#page-12-0); Mittelbach et al. [2007\)](#page-12-0), and to the higher survival rate (hence greater age) of tropical lineages (Hawkins et al. [2006](#page-12-0); Ricklefs and Renner [1994;](#page-13-0) Weir and Schluter [2007\)](#page-13-0). This implies that rather than to environmental adaptation, phenotypic evolution in tropical areas is usually attributed to the long-term effects of biological interactions (Van Valen [1973;](#page-13-0) Stenseth [1984\)](#page-13-0). Recent studies are now challenging this conventional wisdom. One of the most compelling evidence for this concerns the richest Neotropical ecosystem of them all: the Amazonia (Bush and de Oliveira [2006](#page-11-0); Hoorn et al. [2010\)](#page-12-0). The high vertebrate richness in the Amazon basin traces back to the Miocene (ca. 20 Ma) and appears to be linked to a large-scale geological event (i.e. the beginning of the Andean uplift) that modified the South American landscapes since (Hoorn et al. [2010](#page-12-0)), rather than to the great age of the Amazon rainforest (which traces back to the Eocene, Burnham and Johnson [2004](#page-11-0)).

Recent phylogeographic studies on Amazon mammals provide strong support for the influence of pre-historical environmental changes in shaping species diversification and distribution. This is apparent in patterns in many New World Primates, such as spider monkeys (genus Ateles, Collins and Dubach [2000](#page-12-0)), howler monkeys (genus Alou-atta, Cortés-Ortiz et al. [2003\)](#page-12-0), squirrel monkeys (genus Saimiri, Chiou et al. [2011](#page-12-0)), and capuchins (Lynch Alfaro et al. [2012a\)](#page-12-0). Since molecular studies provide a strong support for the impact of historical and geographical events on Platyrrhini diversification, we might also expect a strong association between geography and phenotypic differentiation, since phenotypic and taxonomic diversification are often associated to each other both cladewise (Rabosky and Adams [2012](#page-13-0)) and in time (Raia et al. [2013](#page-13-0)). This association was in fact reported in at least two groups of New World monkeys: spider monkeys Ateles, whose cranial and dental morphometrics differ across parapatric taxa consistently among environments (Froehlich et al. [1991\)](#page-12-0), and capuchins (Cebinae, genera Cebus and Sapajus) which include allopatric species mostly partitioned into different biomes (Lynch Alfaro et al. [2012a,](#page-12-0) [b](#page-12-0)). Although these studies support a strong impact of geography on phenotypic variation, they do not take environmental variables explicitly into account. Recent evidences and techniques on how environmental variables can be accounted for in ecogeographical studies primarily come from Old World monkeys (Cardini et al. [2007\)](#page-12-0). Cardini and Elton [\(2009](#page-11-0)) investigated skull shape variation in a range of different primates (vervet monkeys and Colobinae) to demonstrate that shape variance within and between species are correlated to the environment (quantified as e.g. annual temperature, precipitation). It is now apparent that environmental variables are a significant factor in explaining skull shape variation in monkeys, although differently among different primate lineages. Indeed, in vervet monkeys environmental shape variance is much higher than in colobines (Cardini et al. [2007;](#page-12-0) Cardini and Elton [2009](#page-11-0)).

Since the geographic range of several South American monkeys spans over highly productive territories, such as the Amazon rainforest, to arid, little productive areas like the Caatinga desert, we expect ecogeographical trends and clines in these monkeys, just as they were identified in their African counterparts.

Here, we test for the association between the environment and skull shape at the macroevolutionary scale in two clades of New World monkeys: howlers (Alouatta spp.) and capuchins (Cebus and Sapajus spp.). These two groups are very different in terms of body mass and feeding ecology. Alouatta species weigh up to 7–8 kg, and are obliged folivore– fruigivore. Capuchin monkeys are far smaller (between 2 and 3 kg), have a frugivorous-omnivorous diet (Rosenberger [1992](#page-13-0)), and most notably use tools (Moura and Lee [2004](#page-13-0)). Whereas they are worlds apart in terms of feeding ecology and size, capuchin and howler monkeys cannot be any more similar in terms of geographic distribution. Species in both genera cover (collectively) most of the Neotropical South America (Cortés-Ortiz et al. [2003;](#page-12-0) Lynch Alfaro et al. [2012a\)](#page-12-0), exhibit little geographic overlap with each other, and most species are limited to a single biome. This makes these two clades ideal to test for the effect of climatic variation on the phenotype, because they experience almost identical climatic variation, whereas the potentially confounding effects of dietary convergence and interspecific competition on shape are not an issue.

We employ a combination of geometric morphometrics and comparative methods to test the hypothesis that environmentally-driven skull shape changes occur similarly in these two phylogenetically and ecologically distant Plathyrrini clades.

We compute and compare direction, size and shape of evolutionary vectors in skull shape of the two monkey lineages. These vectors describe phenotypic changes along an environmental gradient (Adams and Collyer [2009](#page-11-0)) that we defined using biomes as discrete ordinal climatic categories.

Since skull shape in Platyrrhini is profoundly influenced by size (the allometric component, Marroig and Cheverud [2004,](#page-12-0) [2005,](#page-12-0) [2010](#page-12-0)) we test also for allometric signal in our skull shape dataset, then we remove it to analyze ''allometry free'' skull shape components and their association with environmental variables.

Additionally, comparative methods are applied to:

- 1. Detect phylogenetic signal on species averaged skull shapes (Klingenberg and Gidazwiski [2010\)](#page-12-0); and
- 2. Test the association between skull shape and environmental variables at the interspecific scale.

Materials and Methods

Sample Size

Our sample includes 535 wild-caught adult specimens belonging to fifteen different species (Table 1, Electronic Supplementary Material): seven *Alouatta* spp., and nine Cebinae (genera Cebus and Sapajus sensu Lynch Alfaro et al. [2012a\)](#page-12-0). We sampled only South American taxa with the exclusion of species whose range expand into Central America (Alouatta palliata and Cebus capuchinus). In this way, we focus our attention on evolutionary processes occurred within clades, that started roughly at the same moment, when the uplift of Andean Cordillera was completed. The two clades are monophyletic on their own, whereas the tree we produced is paraphyletic as it excludes non-howler Atelidae. This is not detrimental to our analyses as we compared the clades to each other.

For each specimen we photographed skulls in ventral view using a standard protocol: each skull was positioned parallel to a digital camera at a 1 meter distance to avoid parallax and possible image distortion. We used the software tpsDig 2 (Rohlf [2010\)](#page-13-0) to build a database of digital pictures and recorded two-dimensional spatial coordinates of twenty-three homologous landmarks. The landmark configuration was chosen to accurately describe ventralview skull features shared by Alouatta and the Cebinae: the rostrum inclusive of incisor raw, muzzle and palate width, thickness and length of the upper canine, premolars and molars; the occipitalis and the zygomatic arch (Fig. [1](#page-3-0)). Digital landmarking was computed by one of us (N.C.) in order to avoid inter-observer error. Additionally, we randomly landmarked twice sixteen representative specimens and observed a landmark error much smaller than 5 %. This indicates that intraobserver error is negligible (cf. Cardini and Tongiorgi [2003;](#page-12-0) Meloro [2011](#page-12-0)).

Additionally, we recorded the geographical coordinates of specimen collection localities. Geographical coordinates, expressed in decimal degrees, were obtained by using Google Earth and specific reference collection (Silva [2001](#page-13-0)). Data were imported into EsriArcGis 10 software. Under GIS, we sampled for each locality its biome code (Olson et al. [2001](#page-13-0)) and bioclimatic variables [\(www.worldclim.org](http://www.worldclim.org)). Olson et al. [\(2001](#page-13-0)) identified 16 biomes to describe categorically environmental complexity of different ecosystems across the earth. The geographic distribution of our sample

Table 1 Sample size of skull order for taxa and biomes of occurrences

Fig. 1 Disposition of 23 landmarks on a skull of Alouatta caraya specimen. Landmarks definitions: 1 Prosthion: antero-inferior point on projection of pre-maxilla between central incisors, 2 posteriormost point of lateral incisor alveolus, 3 anterior-most point of canine alveolus, 4 mesial P1: most mesial point on P1 alveolus, projected onto alveolar margin, 5–9 contact points between adjacent premolars/molars, projected labially onto alveolar margin, 10 anteriormost point on curvature of the zygomatic process, 11 middle point on curvature of the zygomatic process, 12 posterior-most point on curvature of the zygomatic process, 13 posterior extremity of occipital condyle along margin of foramen magnum, 14 opisthion: posteriormost point of foramen magnum, 15 basion: anterior-most point of foramen magnum, 16 most posterior tip of the palatine, 17 posterior tip onto alveolar margin of M3, 18–23 contact points between adjacent pre-molars/molars, projected lingually onto alveolar margin

covers five of Olson's biomes: $1 =$ Tropical and Subtropical Moist Broadleaf Forest, $2 =$ Tropical and Subtropical Dry Broadleaf Forest, $7 =$ Tropical and Subtropical Grasslands, Savannas and Shrublands, $13 =$ Deserts and Xeric shrublands, $14 =$ Mangroves. Because we are interested in understanding evolutionary changes that occurred within and outside the Amazon, we refined Olson classification for biome 1 as split into two other categories: Amazonian and Atlantic Forest (Fiaschi and Pirani [2009](#page-12-0)). Environmental variables were quantitatively described by 19 different bioclimatic indices (Electronic Supplementary Material). These variables were used to perform a discriminant function analysis, in order to identify a possible climatic gradient between the six biomes we identified.

We anticipate that the chosen biomes could be significantly ordered according to temperature seasonality and annual precipitation from biome "1" (Amazonian forest, category 1), followed by "14" (Mangroves, category 2), then Atlantic Forest (category 3), then biome ''7'' (Savanna, category 4), by biome ''2'' (dry forest, category 5) and finally the driest category biome ''13'' (desert, category 6). This ordinal categorization was necessary for further analyses (see Electronic Supplementary Material for more details).

Geometric Morphometrics

Generalised procustes analysis (GPA) was employed to remove from the original landmark coordinates differences in size and orientation (Rohlf and Slice [1990;](#page-13-0) Adams et al. [2004](#page-11-0)). The algorithm applies translation and scaling to unit centroid size (=the square root of the sum of squared distances between each landmark and the centroid, Bookstein [1989](#page-11-0)) after iterative rotation in order to minimize the summed squared Euclidean distances (procustes distances) among specimens, hence allowing to optimize landmarks superimposition among specimens.

After GPA, original landmark coordinates are transformed into shape variables (the procustes coordinates) while size is obtained as centroid size (Rohlf [2000\)](#page-13-0).

The log transformed centroid size was first analyzed to detect differences between clades, species, biomes and their interaction using standard ANOVA. A test of multivariate allometry was employed using MorphoJ (Klingenberg [2011\)](#page-12-0) in order to identify the strength of allometric signal in shape variables (Klingenberg [1996](#page-12-0); Marroig and Cheverud [2005\)](#page-12-0). MANCOVA was applied to test for differences in allometric slope between clades.

Due to the large size differences between Cebinae and Alouatta spp., two alternative approaches were applied before summarizing multivariate dimensionality of shape variables with principal components. First, a variance covariance matrix was generated using within-species covariances (Thorpe [1983](#page-13-0), [1988](#page-13-0); Klingenberg and Spence [1993](#page-12-0)). Marroig and Cheverud ([2001,](#page-12-0) [2004](#page-12-0)) consistently demonstrated that the within-group covariance matrices are broadly similar between New World Primates. Consequently, by applying this procedure the interspecific differences are minimized to better detect any ecogeographical signal in the principal component analysis (PCA). MorphoJ (Klingenberg [2011\)](#page-12-0) was used to generate the within-species covariances and then to perform PCA on the latter.

A second approach to reduce size and clade differences was to generate "size-free" shape variables obtained as the residuals of multivariate allometry. These coordinates were then submitted to a second PCA in order to visualize "allometry free" shape differences. Thin plate spline was used to visualize shape changes described by Principal Component scores.

Skull Shape and Environmental Adaptation

Every skull specimen is ascribed to a biome and clade (Alouatta or Cebinae either) categories. Before applying the procedure to identify the relative impact of environmental adaptation on skull shape, we performed a MANOVA analysis to test for significant differences in skull shape between ''clades'' and between ''biomes'' (with interaction). In spite of the recognized sexual dimorphism in Alouatta and Cebinae (Flores and Casinos [2011\)](#page-12-0) we noted no significant interaction between ''sex'' and either ''clades'' or ''biomes'' in skull shape. Consequently, and to retain the original sample size, we did not divide our sample into males and females.

We first used the test of Adams and Collyer [\(2009](#page-11-0)) to compare vectors of phenotypic transformations across the environmental gradient between the two monophyletic clades: Alouatta versus Cebinae (inclusive of Cebus and Sapajus). Our environmental gradient (or evolutionary level, sensu Adams and Collyer [2009\)](#page-11-0) is a categorical variable that can be commonly identified in the sample of Alouatta and Cebinae. We used biomes as a potential indicator of the environmental complexity (level). The geographic distribution of skull specimens for both clades covers evenly the six tropical biomes (see previous section).

An ''averaged'' skull shape was computed for each biome within each clade. Averaged skull shapes were obtained first on the principal component shape space of the pooled within-species covariance matrix, then on PCA of ''sizefree'' shape variables. These shapes were used to detect evolutionary vectors (Adams and Collyer [2007](#page-11-0)). The vector that connects one ''averaged'' biome shape to the next has three components: direction, size and shape. By employing the algorithm of Adams and Collyer ([2009\)](#page-11-0) we compared these three vectors' attributes between the two clades. If biomes represent an evolutionary level with common impact on skull shape changes in Alouatta and Cebinae, we expect similarities in all of the three vectors' attributes.

Secondly, in order to increase the resolution of environmental complexity, we also tested for the association between skull shape and 19 standardized bioclimatic variables extracted independently from the collection localities' geographical coordinates. Skull shape coordinates were first averaged by localities for each different species to avoid pseudo-replications (cf. Cardini et al. [2007;](#page-12-0) Cardini and Elton [2009](#page-11-0); Cardini et al. [2013\)](#page-11-0). Two block partial least squares was then applied (Rohlf and Corti [2000](#page-13-0)) with block 1 being represented by the whole skull shape variables (within-species covariances matrix and ''size-free'' coordinates), and block 2 by the 19 standardized bioclimatic variables. Here, two block PLS is preferred over linear regression because it does not assume any statistical dependency between variables yet maximizes vector correlation between blocks (Zelditch et al. [2004](#page-13-0)). Similar examples for geometric morphometrics and climatic datasets can be found in Monteiro et al. ([2003\)](#page-13-0), Frost et al. [\(2003](#page-12-0)) and Piras et al. ([2012\)](#page-13-0). Additionally, to test the hypothesis that a common environmental gradient influences skull shape, we first computed PLS vectors independently for the two clades, than compared the vector angle between these two groups to a random distribution (Klingenberg et al. [2002](#page-12-0)) under the null hypothesis of independence. If the observed angles are smaller than expected by chance (at $p < 0.05$) then non-independence occurs between vectors.

For these analyses we used the R code as published in Adams and Collyer [\(2009](#page-11-0)) together with tpsPLS ver. 1.18 (Rohlf [2006\)](#page-13-0) and MorphoJ (Klingenberg [2011\)](#page-12-0) for partial least squares analyses.

Comparative Methods

To validate the results obtained for the whole skull shape sample at the macroevolutionary scale, we re-computed skull shape space based on species averages. Procustes registration was applied separately on each species of Alouatta and Cebinae and a new shape space (based on the within-species covariance matrix, see previous section) generated resulting in a dataset with 15 data points (one for species, cf. Meloro et al. [2008](#page-12-0)). A phylogenetic tree based on the topology and time of divergences obtained inde-pendently by Cortés-Ortiz et al. [\(2003](#page-12-0)) for Alouatta and Lynch Alfaro et al. [\(2012a\)](#page-12-0) for Cebinae (Electronic Supplementary Material) was mapped within the skull shape space following Klingenberg and Gidazwiski ([2010](#page-12-0)). We tested for a significant phylogenetic signal in skull shape by comparing the observed sum of procustes distances between species averaged shapes and reconstructed ancestral node values, and the distribution of these sums obtained by randomizing tip and node values (cf. Klingenberg and Gidazwiski [2010](#page-12-0); Meloro and Jones [2012](#page-12-0)).

Additionally, we tested for skull shape-environmental correlation at the macroevolutionary scale by applying two block PLS on the averaged skull shape sample versus the averaged 19 bioclimatic values (as obtained by the observed sample). If environmental adaptation occurs also across species we might expect a significant association in the reduced sample. Independent contrasts were applied to both shape and bioclimatic variables to repeat PLS in a phylogenetic context (Garland et al. [2003;](#page-12-0) Adams [2008](#page-11-0); Klingenberg [2011](#page-12-0); Bastir et al. [2010](#page-11-0)).

Results

Skull Size

Skull size differ significantly between clades and species (ANOVA $F_{14,520} = 82.559, p < 0.0001$; Fig. [2\)](#page-5-0). Due to heterogeneity of variance (Levene_{14,520} = 6.775, $p < 0.0001$) Dunnett T3 test was applied for post hoc comparisons: all members of *Alouatta* are larger than Cebinae, *Alouatta mac*conelli is larger than Alouatta caraya and Alouatta guariba; within Cebinae Sapajus macrocephalus is larger than Sapajus libidinosus, Sapajus cay (smaller then Sapajus nigritus) and Sapajus apella.

A full ANOVA model shows that significant size differences between specimens from the same biomes (ANOVA

Fig. 2 Box plot of natural log transformed centroid size across different taxa (a) and biomes (b). Black string median, grey box (light Alouatta, dark Cebinae): first interquartile, bar second interquartile

 $F_5 = 3.359$, $p = 0.005$) are due to the interaction with the factor clade (ANOVA $F_5 = 3.432$, $p = 0.005$, Fig. 2). Within Alouatta differences between biomes are significant (ANOVA $F_{5,300} = 2.425$, $p = 0.036$) with specimens from Atlantic forest being smaller than those from the Amazon (after Tukey, Levene_{5,300} = 0.691, $p = 0.631$). In Cebinae size per biome differ as well (ANOVA $F_{5,223} = 4.691$, $p<0.0001$) with specimens from Cerrado being smaller than specimens from the Amazonian and Atlantic forests (after Tukey test, Levene_{5,223} = 9.05, $p = 0.479$).

There is a strong allometric signal in the skull shape of howler and capuchin monkeys (Wilks' $\lambda = 0.0678$, $F_{42,492} = 161.169$, $p < 0.0001$) with size explaining 55.659 % of the sample variance. MANCOVA shows that slope differ significantly between clades (Wilks' $\lambda = 0.654$, $F_{42,490} = 6.176, p < 0.0001$ as well as species (Wilks' $\lambda = 0.142, F_{588,6,229,9} = 1.068, p < 0.0001.$

Skull Shape Variation and Evolutionary Vectors

PCA performed on the pooled within-species covariance matrix allows to summarize the variance of 42 shape variables with the first sixteen vectors explaining together 95.151 % of the total variance.

The plot of the first two PCs shows significant differences in skull shapes of Cebinae and Alouatta are. Along the first PC (49.273 %, Fig. [3a](#page-6-0)) negative scores describe Alouatta skulls with relatively short incisors, small premolar and zygomatic arch area, but large molar area. The occipital foramen is relatively short. The opposite occurs in Cebinae that occupies positive scores of PC1 with enlarged rostrum, incisors and premolars, wider zygomatic arch and occipital foramen (Fig. [3a](#page-6-0)). PC2 (9.93%) is strongly loaded on shape changes in the zygomatic arch area that is shorter and projected more posteriorly in Alouatta (positive scores) while it is larger and projected anteriorly in Cebinae (Fig. [3a](#page-6-0)).

The "size-free" PC scores clearly show that species differences are no longer apparent (Fig. [3](#page-6-0)b) even if Alouatta specimens still tend to occupy positive PC scores with relatively longer rostrum-molar raw and shorter zygomatic arch while Cebinae are in negative scores.

MANOVA reveals significant differences in skull shape between the two clades both when using within-species covariance matrix and ''size-free'' PC scores (Table [2](#page-6-0)). Hotelling pairwise comparisons (based on the first 4 PC scores (=77 %, selected because of sample size restriction) reveal that biomes averages are substantially different between clades (Table [3](#page-7-0)). However, within clades only the specimens from Amazonian forest are significantly different in skull shape from the Atlantic forest and savanna biomes. When using "size-free" PC scores this same conclusion applies to Alouatta only, while in Cebinae also specimens from other biomes (e.g. Mangrove, dry forest) are quite distinct in terms of skull shape.

By plotting averaged skull shapes per biome we observed similarities in vectors of skull shape transformation that are broadly confirmed by statistical tests, no matter if using PC

Fig. 3 Scatter plot of PC1 versus PC2 based on a sample of 535 specimens of Cebinae (circles) and Alouatta (triangles). In a PC scores are from pooled within-species covariance matrix, in b PC scores are plotted after removing allometry. Deformation grids are placed as representatives of the extreme scores of each axis. Large circles/triangles represents averaged phenotype for each biome type (going from black circle biome 1—Amazon forest to white circle biome 6—desert)

Table 2 Hotelling pairwise comparisons

		Pillai trace Approx. $F \cdot df \cdot 1 \cdot df \cdot 2$			\boldsymbol{p}
Pooled within-species					
Clade	0.985	770.67	42	482	< 0.0001
Biomes	0.877	2.46	210	2.430	< 0.0001
C lade \times biomes	0.809	2.23	210	2.430	< 0.0001
Size-free					
Clade	0.345	6.051	42.	482.	< 0.0001
Biomes	0.877	2.46	210	2.430	< 0.0001
C lade \times biomes	0.809	2.23	210	2.430	< 0.0001

scores of within-group covariance or ''size-free'' variables (Fig. 3; Table [4\)](#page-7-0). After 10,000 randomizations, we identified non-significant differences in all of the three vectors attributes for both shape datasets: phenotypic trajectory size $(MD_{1,2})$, direction $(\theta_{1,2})$, and shape (D_{Shape}) are the same in the two clades.

Skull Morphology and Environmental Variation

Non-parametric correlation shows that skull centroid size is significantly correlated to a broad range of bioclimatic variables (Table [5](#page-8-0)). Most bioclimatic variables influence in a similar way size of Alouatta and Cebinae (e.g. BIO8, BIO13, BIO18) even if the latter group correlates with a larger number of bio-climes.

Two block partial least squares identifies a significant correlation between skull shape and bioclimatic variables both when using pooled within-species variance or "sizefree'' across a geographic sample now reduced at 227 data points (Fig. [4a](#page-9-0), b). When using within-species variance only the first pair of singular warps (SW) shows a significant association explaining 73.624 % of the total covariation between the blocks. The correlation coefficient for SW1 shape versus SW1 bioclimate is not particularly high $(r = 0.251)$ and their plot shows only small overlap between the two separate clusters (Alouatta on negative SW1 shape scores and Cebinae on positive SW1 scores, Fig. [4](#page-9-0)a). There is more overlap when ''size-free'' shape scores are analysed (Fig. [4b](#page-9-0)) and the first pair of SW1 explains a much higher percentage of covariation (92.338 %). However their correlation is still weak ($r = 0.250$).

PLS repeated on the geographic subsample of *Alouatta* $(N = 134)$ and Cebinae $(N = 93)$ yields non-significant results when using within-species covariance matrix, but a significant association with "size-free" shape data. Indeed, for Alouatta the first pair of SW1 explains a large percentage of covariance (96.629 %) with a high correlation coefficient ($r = 0.666$, Fig. [4c](#page-9-0)) and the same applies to Cebinae (99.153 % of covariance for SW1, $r = 0.573$, Fig. [4](#page-9-0)d). From the plots it is evident a strong discrimination of Amazonian specimens (positive PLS scores) from all the other biomes (Fig. [4c](#page-9-0), d).

Loading profiles for SW1 bioclimates show that in all the shape datasets analyzed there are very similar patterns with BIO4, BIO12, BIO16 and BIO19 always exhibiting high loadings.

The shape deformations in the overall dataset (Fig. [5a](#page-10-0), b) are strongly influenced by differences between Alouatta and Cebinae with major deformations related to climate described by zygomatic arch and its position relative to the teeth. In Alouatta Amazonian specimens adapted to a less seasonal climate with high and constant rate of rainfall show relatively shorter and thin tooth raw and short zygomatic arch area (Fig. [5c](#page-10-0)), while in Cebinae there is a more posteriorly projected zygomatic area and shorter rostrum—with smaller molar raw (Fig. [5d](#page-10-0)).

The angle between SW1 "size-free" shape vectors of Cebinae and Alouatta is 64.186, which is significantly different from a right angle ($p < 0.001$) indicating that the two vectors are not independent from each other.

Table 4 Summary statistics for differences in phenotypic trajectory size ($MD_{1,2}$), direction ($\theta_{1,2}$), and shape (D_{Shape}) between the clade Cebinae and Alouatta

	Parameter	<i>p</i> value
Pooled within-species		
$MD_{1,2}$	0.0152	0.650
$\theta_{1,2}$	125.46	0.815
D_{Shape}	1.0094	0.077
Size-free		
$MD_{1,2}$	0.0026	0.491
$\theta_{1,2}$	56.834	0.799
$D_{\rm Shape}$	0.996	0.128

Comparative Analyses

The phylomorphospace defined by the 15 species and their ancestral nodes mirrors shape differences observed on the whole sample (Fig. [6](#page-10-0)). Again, we note on the PC1 (92.15 % of total shape variance explained) a clear separation between the clades Alouatta and Cebinae whose distance appears to be proportional to the time of divergence from the tree root. The permutation test evidenced a significant phylogenetic signal in skull shape ($p<0.0001$) after $10,000$ randomizations; observed tree length = 0.0146).

Repeating the partial least squares analysis with skull shape as block 1 and averaged bioclimatic variables as block 2 yields non-significant results for the 15 species data (SW1 = 98.3 % of covariation, $r = 0.421$, $p = 0.177$). PLS loadings for Cebinae and Alouatta are remarkably similar in SW1 bioclimate being correlated strongly with BIO12, BIO19 and BIO18 (in Cebinae) and BIO12, BIO19, BIO17 and BIO16 in *Alouatta* (Table [5](#page-8-0)). The test for angle differences in SW1 shape vector between clades is significantly different from 90 $^{\circ}$ (angle = 51.61, $p < 0.0001$).

PLS repeated on the independent contrasts of 15 species data extracts four significant SW pairs (see Electronic Supplementary Material). SW1 explains a high proportion of covariation between shape and bioclimates (95.99 %) and is again loaded on variables which are much the same as with those observed by analyzing Cebinae and Alouatta separately (BIO12, BIO18 and BIO19, which is basically precipitation and seasonality Table [5\)](#page-8-0).

Discussion

The evolution of capuchins is characterized by an early split between robust (Sapajus) and gracile (Cebus) forms during the late Miocene (Lynch Alfaro et al. [2012b](#page-12-0)). After this major event, capuchins began invading tropical South American habitats, diversifying into different species as

	Alouatta 134 size	Cebinae 93 size	227 pooled	227 size-free	15 size-free	Alouatta 134	Cebus 93	15 pooled	IC 14
BIO1	-0.122	-0.257	0.027	0.009	0.022	0.015	0.014	0.019	0.006
BIO ₂	-0.122	-0.026	0.019	0.039	0.028	-0.026	-0.022	0.024	0.010
BIO ₃	0.116	0.252	0.001	0.013	-0.009	-0.012	-0.005	-0.009	-0.014
BIO ₄	-0.252	0.062	0.806	0.799	-0.298	-0.799	-0.822	-0.317	-0.343
BIO ₅	-0.187	-0.278	0.049	0.028	0.012	0.010	0.005	0.009	-0.005
BIO ₆	0.157	-0.024	0.005	-0.018	0.035	0.029	0.033	0.033	0.022
BIO7	-0.212	-0.072	0.040	0.062	0.019	-0.045	-0.042	0.014	-0.001
BIO ₈	-0.203	-0.230	0.034	0.018	0.019	0.003	0.003	0.015	0.002
BIO ₉	-0.005	-0.250	0.016	-0.003	0.027	0.030	0.027	0.024	0.012
BIO10	-0.173	-0.223	0.036	0.017	0.019	0.007	0.006	0.016	0.003
BIO11	-0.040	-0.246	0.016	-0.002	0.026	0.026	0.026	0.024	0.011
BIO12	0.215	-0.095	-0.475	-0.501	-0.024	0.420	0.401	0.007	0.117
BIO13	0.249	-0.234	-0.070	-0.029	0.046	0.057	0.055	0.006	-0.087
BIO14	0.127	0.375	-0.017	-0.014	0.015	-0.028	-0.010	0.037	0.087
BIO15	0.021	0.021	0.009	0.035	-0.009	-0.016	-0.010	-0.027	-0.076
BIO16	0.156	-0.266	-0.197	-0.127	0.050	0.208	0.181	-0.041	-0.256
BIO17	0.137	0.377	-0.044	-0.076	-0.048	-0.028	0.003	0.032	0.229
BIO18	-0.196	0.189	0.013	0.029	-0.633	-0.170	-0.156	-0.598	-0.448
BIO19	0.274	0.067	-0.268	-0.280	0.705	0.320	0.314	0.730	0.726

Table 5 Loading coefficients on SW1 block variables based on different PLS analyses with shape as block 1 and bioclimate as block 2

Variables and values generally higher than the others are in bold

new and different biomes were colonized (Lynch Alfaro et al. [2012a\)](#page-12-0). Almost the same partitioning of species per biomes occurred in the evolution of Alouatta. Notably, the most recent common ancestors of both howlers and capuchins date to latest Miocene. This means that the two clades have had very similar timelines and geography in the history of their diversification.

In terms of dietary habits, though, howlers and capuchins are very different. This is primarily reflected in skull size that differ considerably between the two clades (Fig. [2](#page-5-0)) and then in their skull shapes (Figs. [3](#page-6-0), 5) with traits evidently related to their different feeding adaptations. A curved rostrum with enlarged incisors and premolars, together with a large zygomatic arch, placed more anteriorly for an efficient bite force, characterize capuchins, which is expected given their omnivorous diet (cf. Rosenberger [1992;](#page-13-0) Wright [2005](#page-13-0); Wright et al. [2009\)](#page-13-0). The opposite applies to Alouatta spp. that are homogenously adapted to eat primarily leaves and for this need enlarged molars, and narrower incisors, together with a short zygomatic area positioned more posteriorly (Rosenberger [1992;](#page-13-0) Rosenberger et al. [2011](#page-13-0)). Such a broad differentiation in shape between the two clades is clearly the result of the long time past their most recent common ancestor, as revealed by the phylomorphospace analysis (Fig. [6\)](#page-10-0).

In spite of their very distinct ecomorphologies, we identified almost identical directions of skull shape changes across biomes and/or bioclimatic variables in the two

clades. Phenotypic trajectory analysis (Adams and Collyer [2009](#page-11-0)) unequivocally supports non-independent pathways for evolutionary changes in shape of Cebinae and Alouatta. This similarity of phenotypic trajectory in New World monkeys is remarkable for we are not aware of any similar results in other studies [e.g. ammonoid cephaloids across time, Monnet et al. ([2011\)](#page-13-0); or damselfishes across different dietary strategies, Frédérich et al. [2013\]](#page-12-0). The similarity in evolutionary vectors in capuchins and howlers supports the idea that the past environmental and geographical variation these clades shared during their diversification set the pace and direction of their morphological change. When both Alouatta and Cebinae clades started to diversify in the late Miocene (ca. 6 Ma) the separation between the Amazonian biome (tropical rainforest) and savanna was already existent (Hoorn et al. [2010](#page-12-0)). This indicates that what today characterizes environmental variation these clades meet was already there when they diversified. It is important to note that the pattern of biome phenotypic changes occurs with or without including allometric component in our analyses. As expected, size represents the major mechanism of phenotypic differentiation between New World Primates species (Marroig and Cheverud [2005](#page-12-0), [2010](#page-12-0)) thus explaining in our sample almost half of phenotypic shape variance. There is a size differentiation also between specimens from different biomes but our analyses support only a strong differentiation of Amazonian species which are generally larger than the others. With ''size-free'' shape

Fig. 4 Scatter plot showing the first pair of singular warp axes extracted by two block PLS with skull shape as block 1 and bioclimates as block 2. In a the pooled within group PLS was performed on averaged geographic sample of 227 specimen with Alouatta as grey triangles and Cebinae as white dots, in **b** same symbols as in a but using "size-free" shape variables, in c PLS for a subsample of 134 "size-free" Alouatta showing in black the Amazonian specimens while in d same analyses but using 93 geographically averaged Cebinae

data (Table [3\)](#page-7-0) there is a wider discrimination between specimens of the two clades from different biomes thus suggesting that the adaptation to distinct environments is subtle and act strongly on size-residual shape variance.

This is confirmed by two block partial least squares that indicates a significant impact of environmental variation on skull shape (allometry free) with very similar patterns being noticeable in both clades (the. Frost et al. [\(2003](#page-12-0)) also identified a strong association between geography and "size-free" shape data in PLS thus suggesting that the pattern we observe could be found also in Old World Primates. Interestingly, the environmental variables mostly correlated with morphology are precipitation and seasonality (Table [5](#page-8-0)). Cardini et al. [\(2007](#page-12-0)) highlighted the importance of considering precipitation to interpret skull shape variation in Old World monkeys as this variable influences fruit/vegetable availability. In South America precipitation has been very often advocated as a good predictor for primate richness (Lehman and Fleagle [2006](#page-12-0)). Recent theories about the use of stone tools in Sapajus also seems to highlight precipitation as responsible for the spatial distribution of this behavior (Ottoni and Izar [2008](#page-13-0)). The skull regions mostly correlated with precipitation (and with climatic seasonality) are the dentition, and the zygomatic arch (Fig. [5](#page-10-0)). Overall, the shape vectors describe changes from more seasonal (lower annual precipitation) to stable (high precipitation) environments. In Cebinae, high seasonality relates to larger zygomatic arch anteriorly placed, and to larger dentition (features typical of the robust Sapajus capable to deal with a variety of other food rather than fruit, Wright et al. [2009](#page-13-0); Lynch Alfaro et al. [2012b](#page-12-0)). In Alouatta spp. seasonal environments influence again zygomatic area and dentition. Recent studies suggest a higher consumption of mature leaves in Southern/Eastern part of South America for A. caraya and A. guariba (Martins [2008\)](#page-12-0). It seems likely that the relatively higher

Fig. 5 Shape deformations related to the first singular warp going from the most extreme negative score $(left)$ to the most positive scores (right). In a deformations are from pooled within group PLS of 227 specimens, in b deformations from PLS analyses of "sizefree'' shape components of 227 specimens, in c deformation from PLS of a subsample of 134 "size-free" Alouatta while in d same but form the analysis of 93 geographically averaged Cebinae

Fig. 6 Phylomorphospace based on species averages (15 data points) for Cebinae and Alouatta. Deformation grids are placed as representatives of the extreme scores of each axis

productivity of the Amazonian rainforest allows both capuchins and howlers to become less constrained in skull ecomorphology in relation to the annual abundance of energetic fruits that do not demand for high biomechanical

performance. We also note that the ecological differentiation between Cebinae and Alouatta does not necessarily point to completely opposite feeding strategies. In fact, fruits tend to be most equally represented in all diets of

PC₂

different *Cebus*, *Sapajus* and *Alouatta* species (Rosenberger [1992;](#page-13-0) Chapman and Fedigan [1990](#page-12-0); Galetti and Pedroni [1994;](#page-12-0) Agostini 2008; Agostini et al. 2010). Such a preference for fruit, together with ecomorphological features generally correlated to feeding on seed, have recently questioned the complete folivory of the howler monkeys that Rosenberger et al. ([2011\)](#page-13-0) renamed as ''semifolivours''. This means that when fruit, as a resource, is available, both howlers and capuchins feed on it. To the South of Amazonia, where fruit is not nearly as abundant as in the rainforest, the consumption of harder food items forces species in both lineages to adapt.

?>It is important to remark once more that allometry is responsible for most shape variation in both capuchins and howler monkeys. Yet, our results also convincingly point to the importance of environmental settings in patterns of shape change in Primates. Despite *Alouatta* spp. and cebids are different in so many regards, they faced the common problem of exploiting hard vegetables (whether they were fruits, leaves, or seeds) as they moved to the more seasonal South. These Primates reacted in very similar ways, with feeding adaptation visible and convergent in the same morphological districts of the skull. The intensity and direction of shape changes are the same in the two groups of species, suggesting that climatic variation was of paramount importance to adaptation in New World Primates. Clearly, capuchins and howlers are only a small fragment of the complex evolutionary Platyrrhini radiation in the New World, and more clades have to be studied to understand if such a pervasive effect of environmental variation of shape differentiation generalizes to New World monkey as a whole.

Conclusion

Macroevolutionary transformations are often advocated as constrained by phylogenetic history, adaptation or biomechanical requirements (Gould [2002\)](#page-12-0). In tropical regions, environmental variation is claimed to be less important than biological interaction in shaping adaptation (Van Valen [1973](#page-13-0); Stenseth [1984\)](#page-13-0). We demonstrate that in spite of a deep morphological difference between capuchin and howler monkeys (as a consequence of their broadly distinct ecologies and lifestyles), environmental variation can be considered as the major factor in ecomorphological differentiation after body size. This validates the well established theory of ecogeographical barrier that supports macroevolutionary differentiation of a broad variety of South American mammals crossing the Amazon forest (e.g. rodents and marsupials De Vivo and Carmignotto [2004](#page-12-0); monkeys of the genus Ateles Froehlich et al. [1991](#page-12-0) and see the crab eating fox, Cerdocyon thous for an example on intraspecific variation, Machado and Hingst-Zaher [2009](#page-12-0)). Out of Amazonia, both Cebinae and Alouatta evolved with similar direction skull shape to deal with a more seasonal, less stable environmental condition.

Acknowledgments We are grateful curators and staff of the Museu Nacional (MNRJ) (J.A. de Oliveira and S.M. Vaz), Museu Paraense Emı´lio Goeldi (MPEG) (S.M. Aguiar and J.S. Silva Jr.), Museu de Zoologia da Universidade de São Paulo (MZUSP) (M. De Vivo and J.G. Barros), Museu de História Natural Capão da Imbuia (MHNCI) (G.M. SiqueiraTebet), Coleção Científica de Mastozoologia da UFPR (DZUP) (F.C. Passos and I.P. Bernardi), and Museu de CiênciasNaturais da Fundação Zoobotânica do Rio Grande do Sul (MCN/FZB) (L.S. Hoffmann) for the kindly authorization and support to specimens access. Senior author (Carlo Meloro) is currently supported by TE-MASAV/DOTTORI DI RICERCA-ESPERTI/26, Francesco Carotenuto by the program FORGIARE V, and Nilton Cáceres by the "Conselho Nacional de Desenvolvimento Científico e Tecnológico" (CNPq) in Brazil (PDE process number 202267/2011-3).

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