Genetic Factors and Orofacial Motor Learning Selectively Influence Variability in Central Sulcus Morphology in Chimpanzees (*Pan troglodytes*)

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Captive chimpanzees (*Pan troglodytes*) have been shown to learn the use of novel attention-getting (AG) sounds to capture the attention of humans as a means of requesting or drawing their attention to a desired object or food. There are significant individual differences in the use of AG sounds by chimpanzees and, here, we examined whether changes in cortical organization of the central sulcus (CS) were associated with AG sound production. MRI scans were collected from 240 chimpanzees, including 122 that reliably produced AG sounds and 118 that did not. For each subject, the depth of CS was quantified along the superior–inferior plane with specific interest in the inferior portion corresponding to the region of the motor cortex where the mouth and orofacial movements are controlled. Results indicated that CS depth in the inferior, but not superior, portion was significantly greater in chimpanzees that reliably produced AG sounds compared with those who did not. Quantitative genetic analyses indicated that overall CS surface area and depth were significantly heritable, particularly in the superior regions, but less so in the inferior and central portions. Further, heritability in CS depth was altered as a function of acquisition of AG sounds. The collective results suggest that learning to produce AG sounds resulted in region-specific cortical reorganization within the inferior portion of the CS, a finding previously undocumented in chimpanzees or any nonhuman primate.

Key words: central sulcus; chimpanzee; heritability; language; orofacial movements; vocal learning

Significance Statement

Recent studies in chimpanzees (*Pan troglodytes*) have shown that some can learn to produce novel sounds by configuring different orofacial movement patterns and these sounds are used in communicatively relevant contexts. Here, we examined the neuromorphological correlates in the production of these sounds in chimpanzees. We show that chimpanzees that have learned to produce these sounds show significant differences in central sulcus (CS) morphology, particularly in the inferior region. We further show that overall CS morphology and regions within the superior portion are significantly heritable, whereas central and inferior portions of the CS are not. The collective findings suggest chimpanzees exhibit cortical plasticity in regions of the brain that were central to the emergence of speech functions in humans.

Introduction

In primate brains, the central sulcus (CS) separates primary motor and somatosensory cortex. Electrical stimulation of the

precentral gyrus elicits different motor movements that are topographically organized along the superior–inferior plane in both human and nonhuman primate brains (Penfield and Boldrey,

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Figure 2. The sequence of postimage processing steps performed from the raw T1-weighted MRI scans (see text for description). The CS (red) is labeled on the 3D cortical graph output.

chimpanzees were being surveyed for their annual physical examinations. Subjects were first immobilized by ketamine (10 mg/kg) or telazol (3–5 mg/kg) and subsequently anesthetized with propofol (40–60 mg/kg/h) following standard procedures at the YNPRC and NCCC facilities. YNPRC subjects were then transported to the MRI facility and NCCC subjects were wheeled to the mobile imaging unit. The subjects remained anesthetized for the duration of the scans, as well as during the time needed to transport them between their home cage and the imaging facility (between 5 and 10 min) or mobile imaging unit (total time ~5 min). Subjects were placed in the scanner chamber in a supine position with their head fitted inside the human-head coil. Scan duration ranged between 40 and 60 min as a function of brain size.

Seventy-eight chimpanzees were scanned using a 3.0 T scanner (Trio; Siemens) at YNPRC. T1-weighted images were collected using a 3D gradient echo sequence (pulse repetition = 2300 ms, echo time = 4.4 ms, number of signals averaged = 3, matrix size = 320×320 , voxel dimension $0.6 \text{ mm} \times 0.6 \text{ mm} \times 0.6 \text{ mm}$). In addition, 142 NCCC chimpanzees were scanned using a 1.5 T Philips machine. T1-weighted images were collected in the transverse plane using a gradient echo protocol (pulse repetition = 19.0 ms, echo time = 8.5 ms, number of signals averaged = 8, and a 256×256 matrix, voxel dimension $0.7 \text{ mm} \times 0.7 \text{ mm} \times 1.2 \text{ mm}$). After completing MRI procedures, the subjects were temporarily housed in a single enclosure for 6–12 h to allow the effects of the anesthesia to wear off, after which time they were returned to their social group. Post-mortem T2-scans were obtained from 44 chimpanzees that had died from natural causes. For the postmortem scanning, either 4.7 or 7 T magnets were used and T2-weighted images were collected in the transverse plane using a gradient echo protocol (voxel dimension $0.67 \text{ mm} \times$

$0.67 \text{ mm} \times 0.67 \text{ mm}$) pulse repetition = 22.0 s, echo time = 78.0 ms, number of signals averaged = 8–12, and a 256×192 matrix reconstructed to 256×256).

Sulci extraction and measurement. The sequence of postimage processing steps performed on the images are shown in Figure 2, *a–h*. The pipeline of processing used to extract the sulci from the raw T1-weighted image derives from a pipeline initially dedicated to the human brain and freely distributed as a BrainVISA toolbox (<http://brainvisa.info>) (Mangin et al., 2004). To account for the differences in chimpanzee anatomy compared with humans, a number of adjustments were performed before the scans were processed using the pipeline procedure within BrainVISA. Specifically, chimpanzee MRI volumes were skull stripped, cropped, and reformatted at 0.7 mm isotropic resolution using ANALYZE 11.0 software and subsequently imported into BrainVISA. The pipeline process of extracting the sulci from the cortex involved a number of steps (Mangin et al., 2004; Fig. 2*a–h*). To align the template brain, the anterior and posterior commissures were manually specified on the MRI at the point where they intersect with the midsagittal slice. The first step was to correct for spatial inhomogeneities in the signal intensity providing a spatially smooth bias field with a stable distribution of tissue intensities (Fig. 2*b*). Next, the analysis of the signal histogram and mathematical morphology were performed using an automatic analysis of the voxel intensities for the entire brain to obtain a binary mask of the brain (Fig. 2*c*). Adjustments were sometimes needed in the histogram process to determine gray and white matter means for chimpanzee brain scans. The mask was then split into the left and right hemispheres and the cerebellum (Fig. 2*d*). A negative mold of the white matter was computed from the split-brain mask. The outside boundary of this mold results

from a 5 mm morphological closing of the masked hemisphere, filling up the folds. The gray/white interface is the inside boundary that preserves deformations and ensures the spherical topology of the mold (Fig. 2e). Finally the mold was skeletonized to detect cortical folding and topological constraints guaranteed that the resulting surfaces would have no holes (Mangin, 2000; Mangin et al., 2004; Fig. 2f,g). Finally, the folds making up the CS in each hemisphere were selected manually (Fig. 1h) using a 3D visualization interface by following standard anatomical landmarks (Bailey et al., 1950; Hopkins et al., 2014b) and the total surface area and average depth were computed for this sulcus in each hemisphere (Fig. 3).

Quantifying CS depth along the superior to inferior plane. The selected CS was meshed using a triangular mesh and the resulting surface was parameterized to create a longitudinal coordinate system (Coulon et al., 2006; Fig. 4a–d). The parameterization process was constrained by the two ventral and dorsal sulcus extremities, automatically detected using the extrema of the first nonzero eigenfunction of the mesh Laplacian (e_1 and e_2 ; Fig. 4b). From these points, a smooth and quasi-isometric (i.e., with minimal metric distortions) coordinate field was extrapolated, which localizes all mesh surface points according to their relative position along the sulcus between the two extremities (Fig. 4c). The coordinate field extends along the length of the CS from the superior/dorsal ($y = 1$) to the inferior/ventral ($y = 100$) ends of the sulcus. Depth was measured at 100 sulcal length positions in a superior-to-inferior progression along the parameterized sulcal mesh surface. Position 1 was adjacent to the interhemispheric fissure and position 100 was adjacent to the Sylvian fissure (Fig. 4c). At each position y along the length, the depth was computed by measuring a geodesic distance (in millimeters) from the brain envelope to the fundus of the sulcus (Fig. 4d).

Data analysis. For each subject and hemisphere, we computed a total surface area and average depth of the CS (Fig. 3). In addition, we computed individual average surface area and average depth by summing the total left and right hemisphere values and dividing by two. For the parameterization analyses, rather than compare and estimate the total additive genetic variance (h^2) scores for each of the 100 points along the sulcus within each hemisphere, we computed average depth values in intervals of five positions. Therefore, we computed 20 depth measures for each hemisphere along the superior-to-inferior plane (1–20). From these measures, we computed the mean depths between the hemispheres at each position as follows: mean depth = $(R + L)/2$, as well as the difference scores in depth at each position as follows: difference = $(R - L)$. For the difference scores, positive values indicate rightward asymmetries and negative values indicated leftward biases. For all analyses, α was set to $p < 0.05$ and *post hoc* analyses, when necessary, were performed using Tukey's honestly significant difference test.

Quantitative genetic analyses. To estimate heritability, we used the software package SOLAR (Almasy and Blangero, 1998), which uses a variance components approach to estimate the polygenic component of variance when considering the entire pedigree (Rogers et al., 2007; Fears et al., 2009; Fears et al., 2011; Hopkins et al., 2014a). h^2 is the proportion of total phenotypic variance that is attributable to all genetic sources. Total phenotypic variance is constrained to a value of 1; therefore, all nongenetic contributions to the phenotype are equal to $1 - h^2$. We used SOLAR in two ways. First, we used SOLAR to determine heritability in the total surface area, average depth, and regional variation in CS depth for the left and right hemisphere. Second, we used the bivariate function within SOLAR to calculate genetic correlations between CS depth measures in the left and right hemispheres along the superior–inferior plane and between depth measures for the left and right hemisphere CS mea-

Figure 3. *a*, 3D rendering from BrainVISA of chimpanzee brain with the CS extracted from the image to illustrate the sulcus dimensions of length and depth along the superior–inferior plane (*b*) and estimated location of different motor functions based on the original studies by Bailey et al. (1950) (*c*).

asures along the superior–inferior plane within the AG[−] and AG⁺ groups. Heritability estimates and genetic correlations at $p < 0.05$ or less were considered significant.

Results

Descriptive data

Because there are very little data available on CS morphology in nonhuman primates, we initially performed some descriptive analyses. Shown in Table 1 is the average overall CS surface area and mean depth for male and female chimpanzees scanned at 1.5, 3, and 7 T. For the overall surface area and mean CS depth, we performed two mixed-model ANCOVAs with sex and rearing history serving as the between-group factors and scanner and age as the covariates. Hemisphere was the repeated measure. For surface area, we found a main effect for sex of $F_{(1,255)} = 8.698$, $p = 0.003$. Not surprisingly, males had significantly larger CS surface areas than females. No other main effects or interactions were significant. However, we did find that the covariate scanner was significant ($F_{(1,259)} = 17.145$, $p = 0.000$). The average surface area for chimpanzees scanned at 1.5 T was smaller than those scanned at 3 and 7 T (Table 1). For the mean CS depth, no significant main effects or interactions were found; however, as was the case with surface area, scanner was a significant covariate ($F_{(1,259)} = 40.946$, $p = 0.000$). The average mean CS depth was lower in chimpanzees scanned at 1.5 T compared with those scanned at 3 and 7 T (Table 1).

For the parameterization data, we performed two mixed-model ANCOVAs with sex serving as the between-group factor and scanner and age as the covariates. For one analysis, the mean CS depth for the 20 regions was the repeated measure and, for the second analysis, the repeated measure was the CS difference score. For the mean CS depth, significant main effects for CS region ($F_{(19,4788)} = 214.25$, $p = 0.000$) and sex ($F_{(1,252)} = 5.729$, $p = 0.017$) were found, as well as a two-way interaction between sex and CS region ($F_{(19,4788)} = 2.241$, $p = 0.002$). The mean CS depths for male and female chimpanzees are shown in Figure 5. First, for both males and females and consistent with previous reports, chimpanzees showed the classically described pattern of CS depth with higher peaks at the inferior and posterior regions

Figure 4. 3D rendering of the CS. *a*, 3D rendering of the CS extracted from the cortical graph and oriented on a horizontal axis where the left end of the sulcus represents the superior end ($y = 0$) and the right the inferior tip ($y = 100$). *b*, Sulcus parameterized to create a normalized x -/ y -coordinate system with the bottom ridge of the sulcal mesh (i.e., the sulcal fundus; $x = 100$) and the top ridge (i.e., at the brain envelope; $x = 1$) representing the x -axis. The end points of the sulcus where these top and bottom ridges join represent the y -axis. *c*, Depth measured from the brain envelope to the fundus of the sulcus along 100 sulcal length positions in a superior-to-inferior mesh surface. *d*, Finally, from the output text file, we can determine the maximum superior point (SP) of the CS by finding the largest depth value between positions 1–50. The maximum inferior point (IP) is represented by the largest depth value found between positions 51 to 100. The pli de passage frontoparietal moyen (PPFM) is the lowest depth value between the SP and IP positions.

Table 1. Descriptive data on CS surface area and mean depth

Scanner	Females	Males	Mean
Surface area, mm ²			
1.5 T	1410.39 (20.80)	1533.05 (32.49)	1417.72 (20.80)
3 T	1595.53 (32.43)	1643.26 (72.77)	1619.40 (39.87)
7 T	1580.10 (53.02)	1527.01 (48.02)	1553.56 (35.71)
Mean	1528.67 (23.7)	1567.77 (31.02)	
Mean CS depth, mm			
1.5 T	8.941 (0.100)	9.191 (0.130)	9.066 (0.083)
3 T	9.466 (0.129)	9.631 (0.290)	9.548 (0.159)
7 T	9.866 (0.211)	9.638 (0.191)	9.752 (0.142)
Mean	9.424 (0.093)	9.487 (0.124)	

and shallower depth in the central portion. Second, with respect to the sex-by-region interaction, *post hoc* analysis indicated that males had larger CS depths at positions 4, 5, 6, 7, 8, and 20. We also found that the covariate scanner was significant ($F_{(2,252)} = 7.553, p = 0.001$). The overall CS depth was lower in chimpanzees scanned at 1.5 T (mean = 12.88, SE = 0.074) compared with those scanned at 3 T (mean = 13.36, SE = 0.104) and 7 T (mean = 13.17, SE = 0.135). For the CS difference scores, no significant main effects or interactions were found.

Heritability in total CS surface area and depth

In the initial analysis, heritability estimates in overall surface area and depth measures for the left and right hemisphere and the average between the two measures were derived using SOLAR. Covariates included sex, age, rearing history, and scanner. The results are shown in Table 2. Significant heritability was found for the overall, left, and right surfaces areas and average depths. Sex and scanner were significant covariates for the surface area and depth measures. As noted above, males had larger surface areas and depths than females and the chimpanzees scanned at 1.5 T had smaller surface areas and depths than chimpanzees scanned at 3.0 T and scanned postmortem. We also performed genetic

Figure 5. Mean depth (in millimeters) averaged from both the left and right hemispheres distributed along the superior–inferior y -axis of the CS for male chimpanzees (solid square) and female chimpanzees (open circle line).

correlations between the overall left and right surface areas and depth measures. For surface area, significant genetic ($\text{RhoG} = 0.96, p = 0.021$) and environmental ($\text{RhoE} = 0.479, p = 0.0002$) correlations were found. Similarly, for mean depth, significant genetic ($\text{RhoG} = 0.98, p = 0.021$) and environmental ($\text{RhoE} = 0.421, p = 0.002$) correlations were found.

Heritability in CS depth from the superior to inferior positions

In the next set of analyses, we computed the heritability estimates for the parameterized depth measures that were computed along

Table 2. Heritability in overall CS surface area and depth with covariates of scanner and sex ($N = 264$)

	h^2	SE	p	Variance
Surface area				
Left	0.374	0.119	0.0001	0.099
Right	0.339	0.117	0.0002	0.088
Mean	0.403	0.118	0.0001	0.102
Mean depth				
Left	0.286	0.148	0.010	0.157
Right	0.183	0.116	0.035	0.137
Mean	0.298	0.140	0.006	0.173

Figure 6. h^2 estimates for positions 1–20 along the superior to inferior plane. Black triangle is the average for the left and right hemisphere, red square is the left hemisphere, and blue circle is the right hemisphere.

the superior–inferior plane. As before, age, sex, scanner, and rearing were covariates in each analysis. The h^2 estimates for each region of the 20 regions for the left and right hemispheres, as well as the average between the two, are shown in Figure 6. For the mean CS depth at each position, significant heritability was found at positions 3 to 8, 15, 16, and 17 (Fig. 6). For the left hemisphere, significant heritability was found at positions 4 to 8 and 20. For the right hemisphere, significant heritability was found at positions 3 to 8 as well as positions 15, 16, and 17. Last, for the difference scores at each CS position, none of the heritability estimates was significant. Therefore, there was no evidence that asymmetries in CS depth were heritable.

Brain–behavior phenotypic associations

In the next analysis, we compared the average CS depth, $(R + L)/2$, and asymmetry in CS depth, $R - L$, at positions 1–20 in relation to sex, gesture handedness, and vocal grouping in two ANCOVAs. For each analysis, CS position was the repeated measure and sex, gesture handedness, and vocal grouping were the between-group factors. Age and scanner were covariates. For the average depths, we found a significant sex-by-region interaction ($F_{(17,3485)} = 2.176, p = 0.003$). *Post hoc* analysis indicated that males had larger mean depth values than females for positions 3–7 and no significant differences were found for the remaining regions.

For the asymmetry measures, we found a significant main effect for gesture handedness ($F_{(2,237)} = 4.117, p = 0.016$) and a

Figure 7. Mean ($\pm$ SE) CS depth asymmetry values for positions 1–20 for AG⁺ and AG[−] chimpanzees. *a*, Males. *b*, Females. On the top is a horizontal view of the CS from the superior (0) to inferior (20) position to illustrate the location of the effect of AG sound use on variation in CS depth asymmetry.

significant three-way interaction between sex, vocal grouping, and region ($F_{(19,4503)} = 1.826, p = 0.015$). For the gesture handedness main effect, *post hoc* analysis revealed that the difference scores were significantly lower in right-handed chimpanzees (mean = -0.122) compared with ambiguously handed (mean = $+0.304$) and left-handed (mean = $+0.166$) chimpanzees. The mean difference scores for each position and vocal group are shown in Figure 7, *a* and *b*, for males and females. *Post hoc* analysis indicated that, in females, difference scores were significantly more leftward in AG⁺ compared with AG[−] individuals at position 17. No other significant differences between AG⁺ and AG[−] individuals were found among females. For males, AG⁺ individuals had significantly greater leftward difference scores at positions 12 and 13 than AG[−] chimpanzees. In addition, at position 17 and, in contrast to females, AG⁺ individuals had significantly greater rightward differences scores than AG[−] chimpanzees. As can be seen in Figure 7, *a* and *b*, the pattern of asymmetry differs substantially between AG⁺ and AG[−] individuals differ from positions 10–19, those regions corresponding to the middle and inferior CS.

Genetic associations between AG sound production and CS depth

In this analysis, we tested whether the phenotypic relationship between AG sound production and CS depth was explained by a common set of genes underlying their expression. For this analysis, using SOLAR, we performed genetic correlations between AG sound production and overall left and right CS surface area and mean depth and the CS depth measures for the left and right hemispheres at positions 4, 5, 6, 7, 8, 9, 16, 17, and 18. These CS positions were selected because the previous analyses found these

Figure 8. h^2 estimates for positions 1–20 along the superior to inferior plane for AG[−] (circles) and AG⁺ (squares) chimpanzees.

measures to be significantly heritable (see above). None of the genetic correlations was significant.

Heritability in CS depths in AG[−] and AG⁺ chimpanzees

Finally, as a means of further examining the role of genetic and experiential factors on variation in CS depth, we performed separate heritability analyses in the AG[−] and AG⁺ chimpanzees. We reasoned that, if learning to produce AG sounds had some consequence on the genetic determination in CS depth either for the left or right hemisphere, then heritability estimates would be higher in AG[−] compared with AG⁺ chimpanzees. The results of these analyses are shown in Figure 8. For the AG[−] chimpanzees, significant heritability in mean CS depths were found at positions 4–11, as well as 16, 17, and 18. In contrast, in the AG⁺ sample, significant heritability was found only at positions 4 and 5. Therefore, heritability in average CS depth was far more robust within the AG[−] compared with the AG⁺ chimpanzees, particularly in the central and inferior regions.

Discussion

Several findings were revealed in this study. First, overall CS surface area and mean depth are significantly heritable in chimpanzees. These results are consistent with previous findings in humans, chimpanzees, and baboons (Kochunov et al., 2005; Rogers et al., 2007; Kochunov et al., 2010; Rogers et al., 2010; McKay et al., 2013; Gómez-Robles et al., 2015) and indicate that the development of the overall CS surface area and depth are influenced by genetic factors. Notwithstanding, our findings also show that, when considering CS depth along the superior to inferior plane, significant heritability within the left and right hemispheres are most pronounced in the regions superior to the plic-de-passage, more spurious in the inferior regions, and absent in the central region. From a comparative perspective, the pattern of heritability and genetic correlations in CS depth along the superior to inferior plane are quite similar to the findings reported in humans by McKay et al. (2013).

A second important finding of this study revealed that chimpanzees who produce AG sounds and those who preferentially gesture with the right hand show larger CS surface areas than those chimpanzees who do not produce AG sounds or who are left-handed or ambiguously handed for manual gestures. The results also indicate that CS depth in the inferior portion of sulcus is greater in the left hemisphere in AG⁺ compared with AG[−] chimpanzees, particularly among males. We believe that the difference in CS depth reported here between the AG⁺ and AG[−] chimpanzees reflect cortical reorganization, particularly in the inferior regions where motor control of orofacial musculature is

represented. Three sets of findings in this report support this conclusion. First, there were no significant genetic correlations between AG sound production and CS depth at those positions that were heritable. Second, recall that heritability in CS depth along the superior to inferior plane was much more robust in AG[−] compared with AG⁺ chimpanzees, suggesting that learning to produce these sounds likely altered the pattern of heritability in CS depth (Fig. 8). Third, the most robust association between AG sound production and CS depth were in regions that were found not to be heritable.

Although the most parsimonious explanation for our findings is that AG sounds are socially learned rather than determined exclusively by genetic factors (Taglialatela et al., 2012), we cannot rule out alternative explanations. Notably, it may be the case that differences in CS depth resulted in some chimpanzees being able to learn to produce AG sounds, whereas others did not. Given the design of this study, we cannot resolve these two different explanations, a problem not unlike the differences in cortical organization often reported in human structure–function comparisons between, for example, musicians and nonmusicians (Gaser and Schlaug, 2003; Li et al., 2010) or under conditions of forced hand use (Klöppel et al., 2010). Notwithstanding this limitation, the results reported here provide the first evidence of an association between orofacial motor control and cortical organization in nonhuman primates and therefore represent an important finding (but see Coudé et al., 2011). Importantly, the collective findings on the neurogenomic correlates of AG sound production provide solid additional support for the notion that their use is in chimpanzees is under voluntary control by the emitters and AG sounds are ultimately related to brain specialization for communication.

With respect to lateralization, we found differences between AG⁺ and AG[−] chimpanzees for the central and inferior CS mean depth and this was particularly the case for males. Previous studies have reported left hemisphere asymmetries in the production of AG sounds in chimpanzees (Losin et al., 2008; Taglialatela et al., 2008; Taglialatela et al., 2011) and the results reported here, at least for mean depth, are consistent with these observations. It is possible that asymmetries in facial expressions associated with the production of AG sounds might better predict asymmetries in the inferior CS depth; however, unfortunately, these data are not available for the subjects in this study. This possibility should be explored in future studies.

Regarding the sex effect of AG sound production and CS asymmetry, it is important to note that a higher proportion of captive male chimpanzees produce AG sounds than females (Taglialatela et al., 2012) and similar findings have been reported for wild individuals (Watts, 2016). Indeed, within the sample of chimpanzees in this study, a significantly higher proportion of males (60%) produced AG sounds compared with females (41%) ($\chi^2_{(1,240)} = 5.297, p = 0.021$). Therefore, the stronger association between AG sound production and CS depth was likely attributable to the fact that this communicative behavior was exhibited more consistently in males compared with females. Recently, Hopkins et al. (2016) reported that the association between measures of brain asymmetry and interhemispheric connectivity also differ between male and female chimpanzees. Males with larger absolute brain asymmetries show reduced interhemispheric connectivity compared with females. Combining these findings with the results reported here suggest that one reason for the reported sex difference in AG sound production and CS asymmetry is the fact that these anatomical regions are simply more lateralized in male compared with female chimpanzees.

One limitation of this study was the fact that we classified chimpanzees into two discrete behavioral categories, AG⁺ and AG[−]. Indeed, it could be argued that AG sound production lies on a continuum instead of a dichotomous scale. Certainly, there are individual differences in the frequency of AG sound production across chimpanzees, but the problem with trying to characterize the distribution on a continuum is that there are many subjects that do not produce any AG sounds. Therefore, there would be a large number of individuals with a score of zero and then some variable distribution of individuals who produce one or more AG sounds within the context of any given experimental paradigm. For example, in the study by Leavens et al. (2010), the communicative behavior of 110 chimpanzees was examined during a 30 s test when a human was looking directly at the focal subject or a chimpanzee immediately adjacent to the focal chimpanzee. In terms of AG sound production, 63 chimpanzees failed to produce an AG sound in either test condition. For the remaining 47 chimpanzees, AG sound production ranged from 1 to 15 occurrences. From a statistical standpoint, treating AG sound production as a continuous variable would be challenging and would not lend itself to correlational types of analyses due to the skewed nature of the distribution. Therefore, the most straightforward and powerful approach was to simply characterize the subjects as producing AG sounds or not (as we did in this study).

In addition, one variable that we were unable to control for was the duration of time that chimpanzees may have been producing AG sounds. In other words, chimpanzees may have acquired the use of AG sounds at different points in life relative to the time that MRI scans were obtained. If changes in cortical folding and organization are attributable to the acquisition and use of AG sounds (rather than the opposite), then the time since the chimpanzee began using AG sounds would be an important variable to consider. Unfortunately, longitudinal data are not available on the acquisition of AG sounds in these two colonies of chimpanzees, so we cannot address this issue. However, we have found that the classification of subjects as AG⁺ or AG[−] as assessed at two time points separated by at least 7 years showed that 72% of the chimpanzees were classified in the same way. Therefore, many of the AG⁺ subjects reliably produced these sounds for an extended period of time before the acquisition of the MRI scans.

In conclusion, it has been hypothesized that voluntary control of the vocal apparatus and associated orofacial musculature was an important step in the evolution of spoken language (Premack, 2004; Lieberman, 2007; Fitch, 2010). Therefore, those individuals early in Hominin evolution who were able to develop these skills may have enjoyed increased fitness compared with other individuals who were not capable of such flexibility in their communicative behavior. Indeed, data from wild chimpanzees indicate that the use of AG[−]-related sounds such as splutters or “raspberries” has an important social function, including the promotion of social bonds, social proximity, and reduction in tension (Fedurek et al., 2015; Watts, 2016). The facilitation of these interactions would have been particularly advantageous in circumstances when the hands were not available for the purposes of producing gestures used in interspecific communication. This scenario is consistent with the gossip hypothesis and the function of these sounds in the context of grooming (Dunbar, 1996, 1998; Dunbar and Schultz, 2007; Leavens et al., 2014). In short, selection for increasing prosocial and cooperative behaviors within the context of grooming may have simultaneously selected for increasing motor control of the vocal and orofacial musculature needed for speech. These collective selective pressures may have

been a driving force for expansion in cortical organization and gyrification within the human brain after the split from the common ancestor with chimpanzees.

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