Morphological Changes of Amygdala in Turner Syndrome Patients

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SUMMARY
Aims: Turner’s syndrome (TS) loses one of the X chromosomes and exhibits social cognition deficits. Previous studies have reported that women with TS demonstrated structural and functional abnormalities in brain, including increased volume in amygdala. However, most studies regarded the amygdala as a whole, and the abnormalities in the specific subregions of amygdala in TS have not been studied. Here, we aimed to investigate the local morphological changes of amygdala in TS using the surface morphology analysis method.

Methods: A total of 19 adolescents with 45X0 TS and 20 matched adolescents with typical development were evaluated using magnetic resonance imaging. The amygdalae of all participants were manually delineated. 3D surface remodeling and parameterization were performed based on the outlined boundaries of amygdalae. We extracted two surface metrics, namely direct Euclidean displacement and normal displacement that were used to represent the morphology of amygdala.

Results: Statistical analysis showed significant outward deformation in the laterobasal subregion of left amygdala in patients with TS, compared with the controls using either direct Euclidean displacement or normal displacement.

Conclusions: Our findings provide novel insight into the pathological changes in the amygdala of patients with TS.

Introduction
Turner’s syndrome (TS) is a common sex chromosome disorder, with the absence of a complete or partial X chromosome in female. Patients with TS have typical physical abnormalities, such as short stature, secondary sexual characteristics dysplasia, and congenital lymphedema [1]. In addition, recent studies have reported the abnormalities in social cognition, involving in facial impressions recognition, emotional stimuli processing and eye gaze perception in women with TS [2–4]. Amygdala plays a major role in emotional processing, for instance fear [5], angry [6,7], and threatening situation [8]. It is also considered to be related with social cognition, particularly emotion recognition and gaze perception [9]. Previous neuroimaging studies have found the enlarged amygdala in patients with TS compared with controls [10–12]. For instance, using volumetric analysis of structural magnetic resonance imaging, researchers found bilateral larger amygdala of women with TS than controls [10,11]. Kesler and colleagues reported the increased volume in left amygdala in patients with TS compared with controls [12]. However, the amygdala is not a homogeneous structure, and it can be divided into three subregions, that is, laterobasal group (LB), centromedial group (CM), and superficial group (SF), according to a histological atlas [13]. Compared with the traditional volumetric analysis method, surface morphology analysis can reveal subtle deformations on the surface of anatomical objects by shape modeling approaches [10]. A number of studies have explored the surface morphology of cerebral structures, such as hippocampus [11,14,15], the central sulcus [16], and amygdala [17–21]. Surface morphology analysis could detect subtle differences on the surface of various brain structures across subjects under neurological disorders. For example, Tamburo and colleagues captured no differences in amygdala volume between the patients with late-life depression and controls, but significant differences in the subfield shape of amygdala by the method of surface analysis [22]. Similarly, Posener and colleagues found highly significant shape differences on the surface of hippocam-
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Materials and Methods

Participants

The subjects included 19 adolescents with 45XO TS and 20 adolescents with typical development, matched for ages (as showed in Table 1). All patients with TS were monosomic females with a 45XO karyotype, who were recruited from the China-Japan Friendship Hospital (CJFH) and Peking Union Medical College Hospital (PUMCH). Control subjects were recruited through local community and parent networks. For each participant, the cognitive assessments were performed within 2 days prior to or after the MRI scan. The participants were assessed with the Chinese version of the Wechsler Intelligence Scale for Children-Fourth Edition (WISC-IV) [25]. The demographic data of all participants are demonstrated in Table 1. All participants had no history of neurological and psychiatric diseases. The research protocol was approved by the Research Ethics Committee of the Beijing Normal University. For each participant, informed written consent was obtained from the legal guardian.

Image Acquisition and Preprocessing

All subjects were scanned on the same 3.0 T Siemens Tim Trio MRI scanner in the Imaging Center for Brain Research, Beijing Normal University. High-resolution 3D T1-weighted images were acquired through magnetization prepared rapid gradient echo (MPRAGE) sequence with the following parameters: echo time (TE) = 3.39 ms; repetition time (TR) = 2530 ms; inversion time (TI) = 1100 ms; matrix = 256 × 256; in-plane resolution = 1 × 1 mm; 176 sagittal slices; thickness = 1 mm; and FA = 7°. All original MR images were registered into stereotaxic space by a linear transformation firstly [26,27]. At the same time, we used the N3 algorithms to correct nonuniformity artifacts for all images [28].

Manual Delineation of Amygdala

Two raters were blind to diagnosis when tracing the amygdala boundary in coronal slices using itk-SNAP (http://www.itk-snap.org) software. This software allows raters to examine the three orthogonal planes together. The amygdala was extracted according to the anatomical definition of amygdala that was described by the Center for Interdisciplinary Brain Sciences Research (CIBSR) at the Stanford University School of Medicine (http://cibsr.stanford.edu) [29]. We used the same criteria with the previous literature, which has the detailed descriptions on the definition of amygdala boundary. Briefly, the most anterior slice of the amygdala was located where anterior commissure (AC) was clearest (thickest, longest, and most continuous). The inferior boundary was formed by a white matter tract at the bottom of the amygdala (as show in Figure 1A). The medial border was always marked by white matter or cerebrospinal fluid (CSF). The superior border of the amygdala was delineated by either high-intensity white matter or CSF (as showed in Figure 1B). Moreover, the lateral border of the amygdala was defined by dense, central white matter tract of the temporal lobe. It was challenging that the amygdala and hippocampus both appeared in the transition (jumping) slice. On these slices, the temporal horn started to enlarge more superiorly along the lateral side of the two structures, and we could divide the two structures by a white matter tract or CSF (as showed in Figure 1C). Raters must examine these slices carefully and discern the amygdala and hippocampus accurately, in the meanwhile, referring to the sagittal view [13] (as showed in Figure 1D).

It is necessary to check the reliability of the manual outlining in order to eliminate the influence of raters’ subjectivity. Two raters who were blind to the side of the brain (left or right) traced the boundary of amygdala on five randomly selected brain volumes. Correlation coefficients between the raters were, respectively, 0.91 and 0.90 for the volumes of left and right amygdala. Six months later, one rater repeated to trace the boundary of amygdala on five randomly selected brains, and the intrarater correlation coefficients for the left and right amygdala were 0.97 and 0.96, respectively.

Surface Modeling and Registration

We utilized the software of SPHARM Modeling and Analysis Toolkit (SPHARM-MAT) (http://www.iupui.edu/~shenlab) which created parametric surface models using spherical harmonics to perform the surface reconstruction of the amygdala [30]. The spherical harmonics description method was a powerful shape descriptor to pinpoint regional changes in morphology. Our surface morphology analysis was based on this method.

As SPHARM-MAT can only be used to model arbitrarily shaped but simply connected 3D objects, each of the manually segmented 3D binary images needed to have a spherical topology before surface reconstruction. Then, the fixed images were parameterized based on spherical mapping, where a con-

<table>
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<th>Table 1 Subject demographics</th>
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<tr>
<td>Turner’s syndrome (n = 19)</td>
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<tr>
<td>Age (years), mean ± SD</td>
</tr>
<tr>
<td>14.0 ± 2.8</td>
</tr>
<tr>
<td>10.2 – 18.6</td>
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<tr>
<td>IQ, mean ± SD</td>
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<td>110.2 ± 14.9</td>
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Continuous and uniform mapping from the object surface to the surface of a unit sphere was created with spherical parameterization. Simultaneously, we used the method of Control of Area and Length Distortions (CALD), which minimized the area distortion cost (ADC) and controlled its worst length distortion cost (LDC) at the same time. Each vertex on the surface was represented by a pair of coordinates after the parameterization. Then, the object surface was expanded into a complete set of spherical harmonic basis functions that were essentially Fourier basis functions defined on the sphere. The Fourier coefficients of the spherical harmonic basis functions that were up to a user-desired degree could be used to reconstruct the object surface while using more coefficients led to a more detailed reconstruction [31].

Surface alignment was an important procedure allowing for pairwise processing or group analysis across different surface models and could facilitate the comparison between the parameter space and object space. This procedure in our experiment mainly included three steps. Firstly, we conducted the first order ellipsoid (FOE) alignment that was performed by selecting the first order spherical harmonic coefficients. Given that the final results of FOE alignments did not preserve the original geometric information in the object space, further alignment was needed. Then, all the intermediate results of the FOE alignment in each side of the amygdala were averaged to generate the corresponding template atlas. Finally, SPHARM Registration with ICP (SHREC) algorithm was used to perform further alignment of the surface of right or left amygdala by minimizing the mean square distances between the corresponding points in the individual surface and template [31].

Surface Metrics

We employed two widely used surface metrics to localize shape changes between individual amygdala and the template. One of the surface metrics was the Euclidean distance from each surface vertex in the subjects to the evenly distributed template [21]. The other surface metric was defined as the distance between each vertex of the subject and the corresponding vertex in the template surface along the normal direction of the mean surface [32]. The positive value of the metrics indicated an outward deformation in the subregion of the subject, relative to the corresponding region of average surface and vice versa. Finally, we used a 5 mm surface-based diffusion smoothing kernel to blur the metric map [33].

Statistical Analysis

For each surface metric, we used a multiple linear regression model to explore the between-group differences vertex by vertex, with age and IQ as covariates. We also evaluated the interaction effects between the group and age. We applied random field theory for multiple comparison corrections on the entire surface [34]. P values <0.05 (FWE-corrected, cluster level) were considered to
be statistically significant. Here, SurfStat (http://www.math.mcgill.ca/keith/surfstat/) was applied to implement the statistical analysis.

Results

Amygdala Volumetry

We computed the number of voxels in the mask of manually segmented amygdala as the volume for left and right amygdala. The volume of the subjects with TS was 1761 ± 214 mm$^3$ for left amygdala and 1810 ± 205 mm$^3$ for right amygdala. And the volume of control group was 1694 ± 217 mm$^3$ for left and right amygdala, respectively. The box plot of amygdala volumes is shown in Figure 2. There were no significant volume differences between the left and right amygdala between the two groups (all $P > 0.05$).

Local Shape Variation

We computed the between-group differences in direct displacement on the surface of amygdala in vertex-by-vertex style. The regions showing significant differences were located in the LB part of the left amygdala. Specifically, the participants with TS showed larger outward deformation, compared with normal controls. The results are illustrated in Figure 3. Figure 3A, B is the scenographs of left amygdala on the coronal plane. The amygdala was displayed from the direction of anterior to posterior. The significant surface cluster ($P < 0.01$, cluster level, corrected) was colored. Figure 3C, D is the back views of Figure 3A, B. Notably, right amygdala did not show any significant differences between the two groups. Using normal displacement, we observed the similar results (see Figure 4). Notably, for each of the surface metrics, we did not find any significant interaction effect between the group and age.

Discussion

In this study, we employed a surface shape analysis method to investigate the abnormal changes of subregional morphology of amygdala in patients with TS. There were no significant differences in bilateral amygdala volumes between the TS and control group. However, we found outward deformations in the LB of the left amygdala surface in the TS group compared with the normal controls. These results provide new insight into the neuropathological basis of TS.

There were some studies reporting the amygdala volume changes in TS groups, and the findings were still inconsistent. For instance, Kesler and colleagues found an increased volume of the left amygdala in 45XO TS compared with the controls [12]. Other studies suggested significant enlargements of bilateral amygdalae for the participants with 45XO TS [10,11]. In contrast, Cutter and colleagues found no significant differences in volumes of the left and right amygdala between the patients with TS and the controls.
Our findings were in line with the results of Cutter and colleagues. However, we found the outward deformations of the left amygdala surface in TS. One possible reason for this finding is that the local deformation changes in patients with TS may be too subtle to make the change of entire volume notable.

In our study, the observed significant changes in the TS group were mainly located around the LB of the left amygdala, which mainly consists of the lateral and basolateral nucleus. The LB is associated with reception of visual information stimulus including facial expression and gaze direction, and fear response. Converging evidence indicated that the LB could integrate visual information, and had axonal connections with visual areas in monkey studies [36–38]. Recently, in vivo study in human demonstrated that the LB of human amygdala is the hub for visual information input processing [39]. The LB is believed to receive related inputs from the cortical and subcortical regions, and sends outputs to other subregions of amygdala and other brain structures [40]. Skuse and colleagues suggested that the left and right amygdala played distinct, but complementary, roles in fearful emotion recognition [41], and the left amygdala activation was more likely associated with the discrimination of facial expression and cognitive processing [41–43]. Hurlemann and colleagues further demonstrated that the LB of the left amygdala was associated with facial emotion perception in a functional MRI study [42]. Recently, some studies have found aberrant social cognition in women with TS, involving fearful face recognition and eye gaze processing [4,41,43]. Particularly, the changed behaviors were related to the amygdala development. Our findings on outward deformation of the LB in the left amygdala therefore provide further support to the notion that the LB of the left amygdala plays a significant role in fearful emotion recognition and gaze perception.

Some issues need to be considered. First, the cerebral pathological changes in TS would be affected by genomic imprinting. Individuals with TS who preserved the maternal X chromosome (45, Xm) and individuals with TS who preserved the paternal X chromosome (45, Xp) had distinct anatomical structure and psychosocial functioning [44,45]. Thus, it is better to take genomic imprinting into consideration in the future. Second, in this study, we did not include any clinical measures for our subjects. It therefore limited our ability to determine the clinical relevance of our findings on structural alterations of amygdala. Third, our study contained a small sample size of subjects, resulting in a limited statistical power. Fourth, we used manual segmentation to outline the boundaries of amygdala, but it is time-consuming and heavy workload. In future, it would be interesting to compare the results using manual segmentation and automatic segmentation methods, such as FSL/FIRST or Freesurfer software (http://surfer.nmr.mgh.harvard.edu/).

**Conclusion**

In this study, we found surface outward deformation of the LB in the left amygdala in adolescents with TS, compared with healthy controls. This particular abnormalities may underlie impaired social cognition in TS, including fearful facial impressions recognition and eye gaze perception. This findings may be useful to enhance our understanding about the pathological changes of the amygdala in patients with TS.

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**Conflict of Interest**

The authors declare no conflict of interest.
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