A voxel-based morphometry study of grey matter loss in fragile X-associated tremor/ataxia syndrome

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Fragile X-associated tremor/ataxia syndrome is a neurodegenerative disorder that primarily affects older male premutation carriers of the fragile X mental retardation gene. Although its core symptoms are mainly characterized by motor problems such as intention tremor and gait ataxia, cognitive decline and psychiatric problems are also commonly observed. Past radiological and histological approaches have focused on prominent neurodegenerative changes in specific brain structures including the cerebellum and limbic areas. However, quantitative investigations of the regional structural abnormalities have not been performed over the whole brain. In this study, we adopted the voxel-based morphometry method together with regions of interest analysis for the cerebellum to examine the pattern of regional grey matter change in the male premutation carriers with and without fragile X-associated tremor/ataxia syndrome. In a comparison with healthy controls, we found striking grey matter loss of the patients with fragile X-associated tremor/ataxia syndrome in multiple regions over the cortical and subcortical structures. In the cerebellum, the anterior lobe and the superior posterior lobe were profoundly reduced in both vermis and hemispheres. In the cerebral cortex, clusters of highly significant grey matter reduction were found in the extended areas in the medial surface of the brain, including the dorsomedial prefrontal cortex, anterior cingulate cortex and precuneus. The other prominent grey matter loss was found in the lateral prefrontal cortex, orbitofrontal cortex, amygdala and insula. Although the voxel-wise comparison between the asymptomatic premutation group and healthy controls did not reach significant difference, a regions of interest analysis revealed significant grey matter reduction in anterior subregions of the cerebellar vermis and hemisphere in the asymptomatic premutation group. Correlation analyses using behavioural scales of the premutation groups showed significant associations between grey matter loss in the left amygdala and increased levels of obsessive–compulsiveness and depression, and between decreased grey matter in the left inferior frontal cortex and anterior cingulate cortex and poor working memory performance. Furthermore, regression analyses revealed a significant negative effect of CGG repeat size on grey matter density in the dorsomedial frontal regions. A significant negative correlation with the clinical scale for the severity of fragile X-associated tremor/ataxia syndrome was found in a part of the vermis. These observations reveal the anatomical patterns of the neurodegenerative process that underlie the motor, cognitive and psychiatric problems of fragile X-associated tremor/ataxia syndrome, together with incipient structural abnormalities that may occur before the clinical onset of this disease.
Introduction

Abnormalities of the fragile X mental retardation gene (FMR1) are associated with a diverse range of behavioural and clinical phenotypes depending on the type of mutation. Expansions of the CGG trinucleotide repeats in the full mutation range (>200 CGG) are the genetic cause of the FMR1 protein deficiency that underlies the fragile X syndrome (Fu et al., 1991; Pieretti et al., 1991; Verkerk et al., 1991). Smaller expansions of 55–200 repeats are referred to as the premutation. Because the premutation does not cause severe protein (FMR1 protein) deficiency as observed in the full mutation, it was initially thought not to be associated with a particular psychological or neurocognitive phenotype. However, recent studies have accumulated evidence for several cognitive and psychiatric problems in adult and child carriers of premutation alleles (Franke et al., 1998; Johnston et al., 2001; Hagerman and Hagerman, 2002; Moore et al., 2004; Cornish et al., 2005; Farzin et al., 2006; Hessl et al., 2007).

Fragile X-associated tremor/ataxia syndrome (FXTAS) is probably the most clinically significant central nervous system phenotype of the FMR1 premutation. FXTAS is a late-onset neurodegenerative disorder primarily affecting older male premutation carriers. Although it is principally characterized as a movement disorder involving intention tremor and gait ataxia, cognitive decline and psychiatric problems are also commonly observed (Hagerman et al., 2001; Jacquemont et al., 2003; Bacalman et al., 2006; Bourgeois et al., 2009). While its pathogenetic mechanism is still unclear, an RNA toxic ‘gain-of-function’ model has been proposed based on several observations (Hagerman et al., 2001; Hagerman and Hagerman, 2004), including: the presence of elevated FMR1 messenger RNA among premutation carriers without clear indications of abnormal FMR1 protein expression (Tassone et al., 2000; Kenneson et al., 2001); the presence of FMR1 messenger RNA within intranuclear inclusions (Greco et al., 2006), one of the hallmarks of FXTAS (Tassone et al., 2004); and premature neuronal cell death in culture combined with dysregulation of several proteins secondary to elevated messenger RNA (Chen et al., 2010, Garcia-Arocena and Hagerman, 2010).

Previous studies have revealed anatomical abnormalities in several structures of the FXTAS brain. Particularly, abnormalities of the cerebellum have been demonstrated in multiple methodologies. In a clinical MRI study, the middle cerebellar peduncle sign was described as one of the most characteristic neuroradiological features of FXTAS (Brunberg et al., 2002). Cerebellar abnormalities were confirmed by post-mortem histological studies that identified several neuropathological features, including Purkinje cell decreases and spongiform changes (Greco et al., 2002, 2006). Significant loss in whole cerebellar volume has been revealed in the male patients with FXTAS (Cohen et al., 2006), which was further replicated in the female patients in a milder form (Adams et al., 2007). However, there have been no studies that attempted to identify the foci of neurodegeneration within the cerebellum. This point is crucial given the fact that there are multiple functionally different subregions in the cerebellum (Stoodley and Schmahmann, 2009) and therefore structural changes in different subregions may make distinct contributions to motor, cognitive and psychiatric problems in FXTAS. Similarly, regionally selective abnormalities in the cerebral cortex are entirely unclear, although the aforementioned MRI volumetric study revealed significant loss of the whole cerebral volume in the patients with FXTAS (Cohen et al., 2006).

Previous behavioural studies revealed patterns of neurocognitive and psychological deficits of FXTAS (Cornish et al., 2005, 2008; Grigsby et al., 2006, 2007), which provide motivations for examining the possible regionally selective abnormalities in cortical and subcortical structures in the FXTAS brain. One study applied an extensive neuropsychological test battery to male premutation carriers with and without FXTAS and reported that, whereas language and visuospatial/attention functions were relatively spared, patients with FXTAS displayed profound deficits of executive cognitive functions, working memory and declarative verbal memory and learning (Grigsby et al., 2008). Several studies replicated significant deficits of executive functions and working memory not only in FXTAS but also in unaffected premutation carriers (Cornish et al., 2008, 2009). For psychological symptoms, it has been reported that major psychiatric features of FXTAS include increased anxiety, depression, disinhibition and apathy (Berry-Kravis et al., 2007b; Bourgeois et al., 2007, 2009). In a large-scale study examining self-reported psychological symptoms of patients with FXTAS and unaffected premutation carriers, the level of the obsessive–compulsiveness was elevated even in unaffected premutation carriers, whereas psychological symptoms of FXTAS extended into other domains, including anxiety and depression (Hessl et al., 2005). Although one recent region of interest-based volumetric study reported a significant correlation between the right hippocampal volume and the severity of anxiety-related psychological symptoms among female patients with FXTAS (Adams et al., 2010), there has been no study that systematically investigated foci of structural abnormalities that may underlie major neurocognitive and psychological problems in individuals with FXTAS and unaffected premutation carriers.

In this study, we adopted the voxel-based morphometry method to examine the regional grey matter loss in the FMR1 premutation carriers with and without FXTAS. Voxel-based morphometry is an automated analysis for assessment of the regional volumetric change over the whole brain (Ashburner and Friston, 2000). For a set of brain regions whose deficits can be responsible for neurocognitive and psychological deficits of the FMR1 premutation carriers, we performed region of interest analyses to examine associations between the grey matter abnormalities in those regions and the severity of the behavioural problems.
Grey matter abnormalities in FXTAS

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Grey matter abnormalities in FXTAS

(refer to ‘Materials and Methods’ section for selection of regions of interest.) Simple voxel-based regression analyses using either CGG repeat size or level of FMR1 messenger RNA of premutation carriers were also performed to examine the effects of the genetic molecular variables on the grey matter abnormality over the whole brain. Lastly, the same simple regression analysis was applied using a clinical scale for assessment of the FXTAS severity to identify brain regions showing progressive neurodegeneration correlated with the development of FXTAS.

Materials and methods

Participants

We examined the brains of a total of 83 male participants between the ages of 40 and 80 years, 28 healthy control participants, 31 participants with the premutation with FXTAS (PFX+), and 24 participants with the premutation without FXTAS (PFX−). In this study, the premutation range was defined as those with a CGG repeat size of between 55 and 200. CGG repeat size was <45 in all the healthy control cases, so that there was no participant whose CGG repeat was within the ‘grey zone’ (45–54 CGG repeats). Participant demographic information is shown in Table 1. The group of PFX+ was significantly older than the other two groups \( F(2,80) = 5.19, P = 0.008 \). Twenty-six controls, 27 PFX+ and 22 PFX− were assessed for full, verbal and performance-scale IQ using the Wechsler Adult Intelligence Scale (Third Edition). According to one-way analysis of variance (ANOVA), a significant main effect of group was found for performance IQ \( F(3,80) = 5.19, P = 0.019 \), with healthy control and PFX+ participants with FXTAS scores of 0 or 1 were placed in the group, while those with FXTAS scores of 2–5 were designated as PFX−. Participants with the premutation were recruited through pedigree analysis of families containing probands with fragile X syndrome. Controls were recruited from the families and the local community through the University of California Davis Medical Centre. Neurological examinations on all healthy control participants were normal, including absence of tremor and ataxia. A signed, written informed consent was obtained according to the Declaration of Helsinki. The protocol was approved by the institutional review board at the University of California, Davis.

Molecular genetic data

Genomic DNA was isolated from peripheral blood lymphocytes using standard methods (Puregene® Kit; Gentra Inc.). For Southern blot analysis, 5–10 \( \mu \)g of isolated DNA was digested with EcoRi and NruI. Hybridization was performed using the FMR1 genomic digoxigenin-labelled 51812.3 probe. Genomic DNA was also amplified by polymerase chain reaction using primers ‘c’ and ‘f’ (Fu et al., 1991). Hybridization was performed with a digoxigenin-end-labelled oligonucleotide probe (CGG)\(^{10}\). Analysis and calculation of the repeat size for both Southern blot and polymerase chain reaction analysis were carried out using an Alpha Innotech FluorChem 8800 Image Detection System (Tassone et al., 2008).

Total cellular RNA was purified from 3–5 ml of peripheral blood using standard methods (Purescript\(^\text{TM}\) kits, Gentra Inc.; Trizol\(^\text{TM}\), BRL). All quantification of FMR1 messenger RNA were performed using a 7900 Sequence detector (PE Biosystems) as previously described (Tassone et al., 2000).

Assessment of clinical severity of fragile X-associated tremor/ataxia syndrome

For participants with CGG repeat count within the premutation range, a trained physician (RJH) scored the severity of FXTAS on a scale ranging from 0–6 as described by our previous studies (Bacalman et al., 2006; Adams et al., 2007). This seven-point staging scale measures functional impairment as follows: 0 = normal functioning; 1 = subtle or questionable tremor or balance problems with no interference in activities of daily living; 2 = minor but clear tremor or balance problems producing minor interference with activities of daily living; 3 = moderate tremor or balance problems with at least occasional falls and significant interference in activities of daily living; 4 = severe tremor or balance problems requiring the use of a cane or walker; 5 = use of a wheelchair on a daily basis and 6 = bedridden. Premutation carriers with FXTAS scores of 0 or 1 were placed in the PFX− group, while those with FXTAS scores of 2–5 were designated as PFX+, meeting clinical criteria for the diagnosis of FXTAS established initially by Jacquemont et al. (2003).

Clinical data acquisition

To examine the relationship between grey matter loss in specific brain regions and behavioural problems of the FMR1 premutation carriers, PFX+ and PFX− individuals were administered a series of examinations for assessing their psychological and cognitive functioning.

Psychological assessment

The Symptom Checklist-90-Revised (SCL-90-R), a standardized self report inventory of current psychological symptoms (Derogatis, 1994), was used for assessing the severity of psychological symptoms

Table 1 Statistics on participant demographic data

<table>
<thead>
<tr>
<th>Healthy controls (n = 28)</th>
<th>Premutation with FXTAS (PFX+) (n = 31)</th>
<th>Premutation without FXTAS (PFX−) (n = 24)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mean (SD)</td>
<td>Range</td>
<td>Mean (SD)</td>
</tr>
<tr>
<td>Age (years)</td>
<td>58.2 (11.1)</td>
<td>40–79</td>
</tr>
<tr>
<td>Full scale IQ</td>
<td>118.6 (16.7)</td>
<td>84–148</td>
</tr>
<tr>
<td>Performance IQ</td>
<td>116.5 (14.3)</td>
<td>89–144</td>
</tr>
<tr>
<td>Verbal IQ</td>
<td>117.0 (17.3)</td>
<td>76–148</td>
</tr>
<tr>
<td>FXTAS score</td>
<td>NA</td>
<td>2.97 (0.98)</td>
</tr>
<tr>
<td>CGG repeat</td>
<td>28.6 (4.37)</td>
<td>17–34</td>
</tr>
<tr>
<td>FMR1 messenger RNA</td>
<td>1.29 (0.29)</td>
<td>0.63–1.85</td>
</tr>
</tbody>
</table>

NA = data not available; SD = standard deviation.
in premutation participants. In this instrument, 90 items are clustered into the symptom dimensions of somatization, obsessive–compulsive, interpersonal sensitivity, depression, anxiety, hostility, phobic anxiety, paranoid ideation and psychoticism. Among these dimensions, we selected obsessive–compulsive, depression and anxiety as particularly relevant dimensions to psychiatric problems of premutation carriers based on our previous study using the SCL-90-R that revealed high scores for obsessive–compulsiveness (Hessl et al., 2005), and our previous meta-analysis finding of the elevated level of anxiety and depression among premutation carriers (Bourgeois et al., 2009).

We obtained scores in 26 PFX+ cases and 17 PFX− cases as follows: obsessive–compulsive: 64.71 ± 11.27 (mean ± standard deviation) in PFX+ and 55.76 ± 7.88 in PFX−; depression: 63.21 ± 13.16 in PFX+ and 54.79 ± 9.46 in PFX−; anxiety: 58.38 ± 12.10 in PFX+ and 51.23 ± 8.27 in PFX−. PFX+ showed significantly higher scores in the three dimensions than PFX− (obsessive–compulsive: \( t_{41} = 2.843, P = 0.007 \); depression: \( t = 2.275, P = 0.028 \); anxiety: \( t = 2.128, P = 0.039 \)).

Cognitive assessment

We assessed the executive functions and working memory in premutation carriers based on previous behavioural studies. We used the *Behavioural Dyscontrol Scale 2* as a measure of executive cognitive functioning. The Behavioural Dyscontrol Scale consists of nine items and measures the capacity for executive cognitive function that addresses self-regulation over voluntary and goal-directed motor behaviours (Kaye et al., 1990). We used the sum of the nine subitem scores as a measure of executive function. As a measure of working memory, we used the sum of working memory subscales (Working Memory Score) of the Wechsler Adult Intelligence Scale. We obtained Behavioural Dyscontrol Scale from 15 PFX+ participants and 18 PFX− participants, and the working memory score from 20 PFX+ participants and 16 PFX− participants. Mean and standard deviations were: Behavioural Dyscontrol Scale = 14.13 ± 5.51 in PFX+ and 20.05 ± 4.09 in PFX−; working memory score = 32.0 ± 7.36 in PFX+ and 34 ± 8.0 in PFX−. PFX+ showed significantly worse Behavioural Dyscontrol Scale (\( t_{34} = 3.538, P = 0.001 \)) whereas there was no significant difference in working memory (\( t_{34} = 0.938, P = 0.355 \)).

Image acquisition

MRI data were acquired on a 1.5T GE Signa Horizon LX NV/I MRI system package (GE Medical Systems, Milwaukee, WI, USA) using a phased array whole-head coil. A high resolution \( T_1 \)-weighted spoiled grass gradient (SPGR) 3D MRI sequence with 124 contiguous horizontal slices (repetition time = 8.7 ms; echo time = 1.8 ms; in-plane resolution = 0.86 × 0.86 mm; slice thickness = 1.3 mm; flip angle = 15°) was administered. During the scan, a custom-built head holder was used to prevent movement.

Voxel-based morphometry analysis

MRI data were processed using Statistical Parametric Mapping software (SPM5) (Wellcome Department of Cognitive Neurology, London, UK) and its ‘VBM5’ toolbox (http://dbm.neuro.unib-jena.de/vbm/vbm5-for-spm5) running on MATLAB version 7.4.0 (The Mathworks, Inc., Natick, MA, USA). Image registration, tissue classification and bias correction were performed under the ‘unified segmentation’ framework (Ashburner and Friston, 2005). In this framework, the first 40 iterations of the initial segmentation estimation are followed by 40 iterations of bias filed correction and finally 20 iterations are made for warping the prior image to the data. This iterated scheme is repeated until no significant changes occur. Standard International Consortium for Brain Mapping grey matter/white matter templates were used for normalization. All images were modulated by correcting for non-linear warping effects and smoothed with a 12 mm full-width at half-maximum smoothing kernel. Grey matter differences between groups were assessed using the general linear model on a voxel-by-voxel basis over the whole brain volume. Because of the significant age difference between groups (Table 1), we included the age of each participant as the covariate of non-interest. Statistical threshold was set at family-wise error rate corrected \( P < 0.05 \) and a spatial extent threshold (\( k \)) of 100 voxels was used for all the contrasts. We first performed the contrast of healthy controls versus PFX+. For the contrast of PFX− versus PFX+, we used an inclusive mask of healthy controls versus PFX− (family-wise error-corrected \( P < 0.05 \), \( k = 100 \)) to increase the statistical power for detecting the intermediate changes that may occur in PFX−.

Region of interest analysis for cerebellar subregions

Given past neuropathological and neuroradiological findings (Brunberg et al., 2002; Greco et al., 2002, 2006), significant grey matter loss in the cerebellum is expected for the patients with FXTAS. In order to characterize abnormalities within the cerebellum in detail, we parcelated the cerebellum into subregions using the WFU PickAtlas (Maldjian et al., 2003) in which the cerebellum is divided into nine subregions in each hemisphere (lobule III, IV/V, VI, VII, VIII, IX, X, Crus I and Crus II) and eight subregions in the vermis (vermis I/II, III, IV/V, VI, VII, VIII, IX and X) (Schmahmann et al., 1999; Lasek et al., 2006). Using the modulated individual grey matter image, we calculated the mean grey matter density of voxels in each subregion. Because of the significant age difference among groups, the individual mean grey matter value was adjusted for the effect of age by calculating the residual after a linear fit of age. The residual value of each subregion in each participant was fed into a two-way ANOVA of Group and Subregion.

Correlation analysis using neurocognitive and psychological measures

A number of past studies replicated crucial involvement of the lateral and medial prefrontal cortex in executive cognitive processes as well as working memory (Buckner and Koutstaal, 1998; Cabeza and Nyberg, 2000). A recent meta-analysis of functional imaging studies of the cerebellum showed that areas in the Crus I and lobule VI of the hemisphere are most reliably activated in tasks for executive processes and working memory (Stoodley and Schmahmann, 2009). It turned out that, in the comparison between healthy controls and PFX−, prominent grey matter loss was identified in the anterior cingulate cortex and left inferior frontal cortex in the cerebral cortex. In the cerebellum, the bilateral Crus I and lobe VI were also profoundly affected (Fig. 1). Therefore, we hypothesized that grey matter loss in these regions may be responsible for impairments of executive functions and working memory in premutation carriers. To test this hypothesis, we calculated correlation between the mean grey matter density of each of these four regions of interest and behavioural measures of the
Behavioural Dyscontrol Scale and the working memory score. To focus on parts of regions of interest where significant atrophy in PFX+ was identified, voxels in each region of interest was defined by the combination of two binary masks: (i) the contrast map of healthy controls versus PFX+ (family-wise error-corrected $P < 0.05$, $k = 100$) and (ii) the WFU PickAtlas (Maldjian et al., 2003). The mean grey matter density of each region of interest was calculated from all voxels in the individual modulated grey matter image that satisfied both binary masks. The WFU PickAtlas mask of each region of interest was generated as follows: left inferior frontal cortex = ‘Frontal_Inf_Oper_L’ + ‘Frontal_Inf_Tri_L’; anterior cingulate cortex = ‘Cingulum_Ant_L’ + ‘Cingulum_Ant_R’; left (right) Crus I and lobule VI = ‘Cerebelum_Crus1_L(R)’ + ‘Cerebellum_6_L(R)’. The individual mean grey matter value was then adjusted for the effect of age by calculating the residual value after a linear fit of age. We used the residual for the calculation of correlation with either Behavioural Dyscontrol Scale or working memory score. To adjust for multiple statistical tests for each scale (four regions of interest/statistical tests per scale), the Benjamini-Hochberg method was implemented, with the false discovery rate set at 5% (Benjamini and Hochberg, 1995).

There has been evidence that anxiety-related symptoms (e.g. obsessive–compulsive disorder and general anxiety) involve abnormalities in the amygdala, insula, anterior cingulate cortex and orbitofrontal cortex (Paulus and Stein, 2006; Etkin and Wager, 2007; Chamberlain et al., 2008). Past studies indicated that neural correlates of depression involve the hippocampus, in addition to the amygdala and anterior cingulate cortex (Soares and Mann, 1997; Sheline, 2000). In the cerebellum, the aforementioned review study indicated that vermal lobule VII is involved in emotional processing by forming the cerebellar-limbic circuitry (Stoodley and Schmahmann, 2009). Because the contrast of healthy controls versus PFX+ revealed prominent grey matter loss in these regions except for the right amygdala and the right hippocampus (Fig. 1), we performed correlation analyses between the mean intensity of each of these regions of interest and subscales of the SCL-90-R (obsessive–compulsive, depression and anxiety). We extracted the mean intensity of each region of interest using the contrast map of healthy controls versus PFX+ and the WFU PickAtlas in the

Figure 1 Significant grey matter reduction of FXTAS (PFX+) identified by comparison either with healthy controls (HC) or with unaffected premutation carriers (PFX-/C0) in sagittal views. The contrast of healthy control versus PFX+ is shown in the yellow-red colour scale and the one of PFX-/C0 versus PFX+ is shown in the deep-light blue scale. The numbers at the bottom left corner indicate the $x$-axis coordinates of sagittal sections. The statistical threshold was set at $P < 0.05$ (family-wise error rate-corrected). The spatial extent threshold was set at 100 voxels.
same way as was done in the correlation analysis of executive function and working memory. The WFU PickAtlas mask of each region of interest was generated as follows: left amygdala = ‘Amygdala_L’; anterior cingulate cortex = ‘Cingulum_Ant_L’ + ‘Cingulum_Ant_R’; left (right) insula = ‘Insula_L(R)’; left (right) orbitofrontal cortex = ‘Frontal_Mid_Orb_L(R)’ + ‘Frontal_Inf_Orb_L(R)’; the vermal lobule VII = ‘Vermis_7’. Correlation with obsessive-compulsiveness and anxiety was tested for the left amygdala, anterior cingulate cortex, left and right insula, left and right orbitofrontal cortex, and vermal lobule VII. Correlation with depression was tested for the left amygdala, left hippocampus, anterior cingulate cortex and the vermal lobule VII. The Benjamini-Hochberg method was used for the adjustment of multiple statistical tests for each subscale in the SCL-90-R (seven tests for obsessive-compulsive anxiety, and four tests for depression).

**Regression analysis using FMR1 molecular variables and fragile X-associated tremor/ataxia syndrome severity scale**

In order to examine effects of FMR1 molecular variables on grey matter, we performed a voxel-based simple regression analysis over the whole brain using either the CGG repeat size or FMR1 messenger RNA level. We included the age of each participant as the ‘nuisance’ covariate in the model. Because there is a systematic difference in the distribution of both molecular variables between healthy controls and the two premutation groups (Table 1), we included the data of the two premutation groups only for this analysis in order to avoid contamination of group (categorical) effects. We also performed the simple regression analysis using the clinical scale for assessment of the FXTAS severity to identify areas showing progressive grey matter reduction caused by the development of FXTAS. We included age as the covariate of non-interest to isolate the effect of clinical severity from the one of age. For this analysis, we used the data from premutation participants whose FXTAS score was $\geq 1$. We used the contrast of healthy controls versus PFX$^-$ (family-wise error-corrected $P < 0.05$, $k = 100$) as the inclusive mask in the three simple regression analyses.

**Results**

**Group difference in the whole-brain analysis**

For the comparison between healthy controls and PFX$^+$, we found clusters of significant grey matter reduction of PFX$^+$ in multiple brain regions over the cerebrum, cerebellum and subcortical structures (Fig. 1 and Table 2). Particularly prominent grey matter loss was observed in the cerebellum, dorsomedial frontal and parietal regions, orbitofrontal regions, insula, medial temporal regions and lateral prefrontal regions. Significant grey matter increase in FXTAS was mainly found in the bilateral posterior superior/middle temporal gyrus (Table 2). In the comparison between PFX$^+$ and PFX$^-$, we found significant grey matter reduction of PFX$^+$ relative to PFX$^-$ in parts of the cerebellum, dorsomedial prefrontal cortex, and precuneus (Fig. 1 and Table 2). There were no significant voxels showing PFX$^+ >$ PFX$^-$. No significant voxels were identified in the comparison between healthy controls and PFX$^-$ in either direction.

**Region of interest analyses for cerebellar subregions**

A two-way ANOVA (Group $\times$ Subregion) for the 26 cerebellar subregions revealed a significant main effect of Group [F(2, 80) = 9.427, $P < 0.001$] and interaction effect [F(50, 2000) = 2.918, $P < 0.001$]. Follow-up one-way ANOVA was performed for each subregion using the Benjamini-Hochberg method for the adjustment for the multiple tests. We did not observe significant effects of Group in lobules VIII and IX in the vermis, bilateral lobule VIII nor right lobule X in the hemisphere. All other subregions showed a significant main effect of Group ($P < 0.05$; Table 3). According to a post hoc test (Tukey’s Honestly Significant Difference), there was a significant difference between healthy controls and PFX$^-$ in all of the regions of interest (Table 3). Compared with healthy controls, PFX$^-$ showed significant reduction in lobule I/II of the vermis and in lobule III in the left hemisphere ($P < 0.05$; Fig. 2 and Table 3). Significant differences between PFX$^+$ and PFX$^-$ were found in lobules IV/V, VI and VII in the vermis, and lobules IV/V, VI, Crus I and right Crus II in the hemisphere ($P < 0.05$; Fig. 2 and Table 3).

**Region of interest-based correlation analysis using neurocognitive and psychological measures**

In correlation analysis with Behavioural Dyscontrol Scale, none of the four regions of interest for executive function reached significance using the threshold corrected for multiple comparisons (left inferior frontal cortex: $r = 0.392$, $P = 0.0968$; anterior cingulate cortex: $r = 0.300$, $P = 0.119$; left Crus I/lobule VI: $r = 0.265$, $P = 0.1364$; right Crus I/lobule VI: $r = 0.373$, $P = 0.065$), although left inferior frontal cortex and the right Crus I/lobule VI showed significant correlations at an uncorrected threshold ($P = 0.024$ and $P = 0.032$, respectively). In correlation analysis using the working memory score, we found significant correlations in the anterior cingulate cortex ($r = 0.498$, $P = 0.004$) and the left inferior frontal cortex ($r = 0.518$, $P = 0.005$) at the corrected threshold (Fig. 3). The two cerebellar regions of interest, by contrast, showed no significant correlation (left Crus I/lobule VI: $r = 0.192$, $P = 0.263$; right Crus I/lobule VI: $r = 0.289$, $P = 0.117$) with Behavioural Dyscontrol Scale.

In correlation analysis using the score of the obsessive-compulsive symptom dimension in SCL-90-R, only the left amygdala reached the significant level ($P = 0.0126$; Fig. 3). Although there were several regions of interest for which a significant correlation was observed at the uncorrected threshold (anterior cingulate cortex: $r = -0.358$, $P = 0.018$; left insula: $r = -0.316$, $P = 0.039$; right insula: $r = -0.311$, $P = 0.042$), these regions of interest did not reach significant level after correction (anterior cingulate cortex: $P = 0.064$; left insula: $P = 0.091$; right insula: $P = 0.074$). No other regions of interest showed significant correlation (vermis lobule VII: $r = -0.205$, $P = 0.188$; left orbitofrontal cortex: $r = -0.272$, ..., $P = 0.091$; right insula: $P = 0.074$). No other regions of interest showed significant correlation (vermis lobule VII: $r = -0.205$, $P = 0.188$; left orbitofrontal cortex: $r = -0.272$, $P = 0.091$; right insula: $P = 0.074$).
$P = 0.091$; right orbitofrontal cortex: $r = -0.277$, $P = 0.101$.

In the correlation using the depression symptom dimension, only the left amygdala was significantly correlated using the corrected threshold ($P = 0.019$; Fig. 3). No other regions of interest showed significant correlation after correction (anterior cingulate cortex: $r = -0.264$, $P = 0.116$; left hippocampus: $r = -0.301$, $P = 0.100$; vermis VI: $r = -0.093$, $P = 0.554$), although the left hippocampus showed a marginally significant correlation at the uncorrected threshold ($P = 0.050$). There was no region of interest that showed a significant correlation with anxiety after

### Table 2 Significant grey matter differences between groups

<table>
<thead>
<tr>
<th>Region</th>
<th>Cluster size</th>
<th>$x$</th>
<th>$y$</th>
<th>$z$</th>
<th>$z_{max}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Healthy controls &gt; PFX$^+$</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Left and right cerebellar hemispheres, vermis, ventral medial temporal regions</td>
<td>53025</td>
<td>5</td>
<td>-77</td>
<td>-21</td>
<td>6.43</td>
</tr>
<tr>
<td>Cingulate cortex, dorsomedial prefrontal cortex, pre-SMA, SMA, precuneus</td>
<td>22729</td>
<td>-1</td>
<td>40</td>
<td>29</td>
<td>5.95</td>
</tr>
<tr>
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<td>-45</td>
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<tr>
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<td>Bilateral gyrus rectus</td>
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<td>-77</td>
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<td>5.22</td>
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<tr>
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<td>35</td>
<td>-69</td>
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<td>Left angular gyrus</td>
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<td>-60</td>
<td>41</td>
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<td>Right angular gyrus</td>
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<td>62</td>
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<td>37</td>
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</tr>
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<tr>
<td>Left post-central gyrus</td>
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</tr>
<tr>
<td>PFX$^+$ &gt; Healthy controls</td>
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<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Left superior/middle temporal gyrus</td>
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<tr>
<td>275</td>
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<td>-9</td>
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<td>4.78</td>
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<td>10</td>
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<tr>
<td>Left lateral occipital gyrus</td>
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<td>-85</td>
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</tr>
<tr>
<td>135</td>
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<td>-73</td>
<td>14</td>
<td>4.74</td>
<td></td>
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<tr>
<td>Left medial orbital gyrus</td>
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<td>26</td>
<td>-11</td>
<td>4.71</td>
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<tr>
<td>Cerebellar lobules V/VI/VII</td>
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<td>5</td>
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<td>-22</td>
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<tr>
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<tr>
<td>Left cerebellar hemisphere VIIB/Crus I</td>
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<tr>
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<td>135</td>
<td>-5</td>
<td>-31</td>
<td>47</td>
<td>4.08</td>
</tr>
</tbody>
</table>

$PFX^+$ > PFX$^-$
- No significant clusters

Healthy controls > PFX$^-$
- No significant clusters

PFX$^+$ > Healthy controls
- No significant clusters

SMA = supplementary motor area.
controls and PFX/C0 messenger RNA. For the analysis involving CGG repeat using FMR1 molecular variables and Whole brain regression analyses r correction (left amygdala: \( r = -0.252, P = 0.717 \); anterior cingulate cortex: \( r = -0.241, P = 0.421 \); left insula: \( r = -0.217, P = 0.284 \); right insula: \( r = -0.212, P = 0.242 \); left orbitofrontal cortex: \( r = -0.222, P = 0.355 \); right orbitofrontal cortex: \( r = -0.208, P = 0.210 \); vermis VII: \( r = -0.154, P = 0.324 \)).

**Whole brain regression analyses using FMR1 molecular variables and clinical scale of fragile X-associated tremor/ataxia syndrome**

We performed regression analyses using CGG repeat size and FMR1 messenger RNA. For the analysis involving CGG repeat size, we found significant negative effects of CGG repeat size in several clusters in the dorsal medial regions including the supplementary motor area and the dorsomedial prefrontal cortex (Fig. 4 and Table 4). No voxels showed a significant positive effect of CGG repeat size. We did not observe a significant correlation with FMR1 messenger RNA in either direction. In the analysis using the clinical scales of the FXTAS severity, we found significant negative correlations in cerebellar lobule I/VII and the orbitofrontal cortex (Fig. 5 and Table 4). No significant positive correlations were observed.

**Discussion**

Brain abnormalities in PFX+ has been demonstrated in previous MRI volumetric studies based on gross anatomical parcellation as
well as clinical MRI investigations (Brunberg et al., 2002; Cohen et al., 2006; Adams et al., 2007, 2010). However, systematic investigations of possible regionally selective abnormalities have not been performed. To our knowledge, this study represents the first demonstration of a spatial pattern of grey matter reduction in the PFX+ brain using voxel-based morphometry. We identified significant grey matter reduction in multiple regions, particularly in the cerebellum, the dorsomedial frontal-parietal regions, the medial temporal regions and the insula. Correlation analysis using the behavioural measurements of the premutation groups indicated that psychological symptoms and working memory deficits of FMR1 premutation carriers are associated with grey matter loss in the left amygdala and in the left inferior frontal cortex and anterior cingulate cortex, respectively. Regression analyses using FMR1 molecular variables showed a significant contribution of CGG repeat size to the grey matter reduction in the dorsomedial frontal regions. Furthermore, progressive grey matter loss correlated with the severity of FXTAS symptomatology was also revealed in a part of the cerebellum and in the orbitofrontal cortex. These findings identify the pattern of anatomical abnormalities in FMR1 premutation carriers that might provide morphological bases for behavioural problems of this population.

Consistent with the past observations, both the whole-brain voxel-based morphometry analysis and the region of interest analyses revealed profound grey matter loss in the cerebellum of PFX+. Significant grey matter reduction was widespread, affecting almost the entire cerebellum except for some subregions in the inferior posterior lobe such as the lobule VIII. In particular, grey matter reductions in the anterior subregions of the vermis and hemispheres were highly significant (Fig. 2). The anterior vermis has been shown to be critical for the regulation of the postural equilibrium while standing (Diener et al., 1989; Ouchi et al., 1999). Severe atrophy in this region of the cerebellum therefore may be directly responsible for gait ataxia, one of the core clinical symptoms of PFX+. We suggest that analysis using behavioural scales for the severity of ataxia, such as the International Cooperative Ataxia Rating Scale, would be necessary to test this hypothesis. It is important to note that significant grey matter reduction was identified in several subregions in the anterior vermis and hemisphere even among PFX+/C0 (Table 3). This observation raises the possibility that degeneration in this region may be the initial pathological process before clinical signs of FXTAS. Significant grey matter loss was also found in several subregions of the posterior vermis. Previous neuropsychological studies described cerebellar cognitive affect syndrome resulted from damage to the posterior lobe (Schmahmann and Sherman, 1998). A recent meta-analysis of functional imaging studies indicated that areas around the vermis lobule VII are recruited with
high probability in tasks of emotional processing (Stoodley and Schmahmann, 2009). Although these findings suggest that significant abnormality of this subregion may be related to psychiatric problems in FXTAS, this possibility was not supported by our correlation analysis that failed to show significant association between the level of depression, anxiety or obsessive–compulsiveness in SCL-90-R and grey matter reduction in this region.

In the cerebellar hemispheres, large grey matter reductions in PFX+ were observed in parts of the neocerebellum, including Crus I, Crus II and lobule VI. According to the aforementioned meta-analysis of cerebellar activation, these neocerebellar regions, particularly the boundary between Crus I and lobule VI, are frequently activated in executive function and working memory tasks (Stoodley and Schmahmann, 2009), which suggests that abnormalities in these cerebellar subregions may contribute to impaired executive and working memory processes in PFX+. However, the present correlation analysis using the Behavioural Dyscontrol Scale and the working memory score did not support this possibility except for the trend-level correlation between the right Crus I/lobule VI volume and the Behavioural Dyscontrol Scale score. Because executive functions comprise various cognitive processes, it is necessary to examine possible relevance with other behavioural tests of executive functions. Compared with these subregions, grey matter reductions in several posterior subregions were less pronounced. In particular, we did not find evidence of significant abnormality of the hemisphere lobule VIII, a subregion that has been shown to be most severely impaired in spinocerebellar ataxia 17 (SCA17), another neurodegenerative disorder caused by a single gene mutation (Lasek et al., 2006). This dissociation is interesting given the phenotypic overlap between this disease and PFX+. Among the posterior subregions, the lobule IX showed significant grey matter loss. This region has been suggested to form the cerebellar node of the ‘default-mode’ network (Habas et al., 2009). Grey matter loss in this region may thus be related to highly significant abnormalities of cortical nodes of this network, such as the dorsomedial prefrontal cortex and precuneus (Fig. 1).

The comparison between the healthy controls and PFX+ revealed profound grey matter atrophy in several cortical and subcortical regions outside of the cerebellum. In particular, a cluster of significant reduction was identified over the extended areas in the medial surface of the brain between the frontal and parietal regions (Fig. 1). This huge cluster comprises multiple regions including the dorsal anterior cingulate/paracingulate cortex, dorsomedial prefrontal cortex, supplementary motor area, middle and posterior paracingulate cortex and precuneus (Table 2). The most prominent grey matter loss was centred in the dorsal anterior cingulate cortex. Previous functional imaging and brain lesion studies have shown that the anterior cingulate cortex is implicated in both cognitive and emotional processing (Bush et al., 2000). In the cognitive domain, activation of this area has been found under executive cognitive and working memory tasks that require
Figure 4  Significant negative effect of the CGG repeat size on the grey matter (GM) density in the FMR1 premutation carriers. (A) Clusters of voxels in the dorsal medial regions showing significant negative correlation with the CGG repeat size. The blue circles indicate the largest cluster in the supplementary motor area in the sagittal, coronal and axial views. (B) The plot of CGG repeat size and the grey matter density at a voxel of the highest t-value in the supplementary motor area (x, y, z = 5, -6, 59). The y-axis represents the residual from a linear regression of age over the modulated grey matter density at this coordinate.

Table 4  Regression analysis using CGG repeat size and FXTAS severity scale

<table>
<thead>
<tr>
<th>Region</th>
<th>Size</th>
<th>x</th>
<th>y</th>
<th>z</th>
<th>z_{max}</th>
</tr>
</thead>
<tbody>
<tr>
<td>CGG repeat size</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Supplementary motor area</td>
<td>283</td>
<td>5</td>
<td>-6</td>
<td>59</td>
<td>4.26</td>
</tr>
<tr>
<td>Mammillary body/anterior ventral hypothalamus</td>
<td>282</td>
<td>3</td>
<td>-5</td>
<td>-11</td>
<td>4.23</td>
</tr>
<tr>
<td>Right orbitofrontal cortex</td>
<td>165</td>
<td>16</td>
<td>69</td>
<td>-9</td>
<td>4.33</td>
</tr>
<tr>
<td>Left dorsomedial prefrontal cortex</td>
<td>144</td>
<td>-4</td>
<td>28</td>
<td>48</td>
<td>4.25</td>
</tr>
<tr>
<td>Right cerebellar hemisphere lobule IX</td>
<td>142</td>
<td>-13</td>
<td>-42</td>
<td>-54</td>
<td>4.35</td>
</tr>
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<td>109</td>
<td>5</td>
<td>50</td>
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<td>FXTAS severity scale</td>
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<td></td>
<td></td>
</tr>
<tr>
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<td>-78</td>
<td>-23</td>
<td>4.66</td>
</tr>
<tr>
<td>Right orbitofrontal cortex</td>
<td>398</td>
<td>16</td>
<td>69</td>
<td>-11</td>
<td>4.63</td>
</tr>
</tbody>
</table>
attention, cognitive efforts and performance monitoring (Smith and Jonides, 1999; Paus, 2001). The anterior cingulate cortex is also a part of the network for decision making and social behaviour by forming reciprocal connectivity with the amygdala, orbitofrontal cortex and ventral striatum (Vogt et al., 1995; Cavada et al., 2000; Beckmann et al., 2009). Within this network, it has been proposed that the anterior cingulate cortex plays a crucial role in selection of action, especially that which involves effort (Rushworth et al., 2007). Therefore, dysfunction of this area might be particularly related to impairments of effortful executive and working memory processes and apathetic symptoms of this disease (Bacalman et al., 2006). The present study lent support for this view by showing a significant correlation between anterior cingulate cortex volume and working memory score (Fig. 3). Further studies are needed to examine whether apathy-related symptoms are associated with structural abnormalities in this region.

The cluster of significant grey matter reduction also extended into the adjacent regions such as the dorsomedial prefrontal cortex, pre-supplementary motor area and supplementary motor area (Fig. 1 and Table 2). The medial dorsal prefrontal cortex and precuneus are crucial nodes of the default-mode network (Gusnard and Raichle, 2001; Raichle et al., 2001). Although the function of this network remains to be clarified, past studies proposed that it is involved in episodic memory retrieval, self-reflection and stream-of-consciousness (Greicius et al., 2003; Cavanna and Trimble, 2006). Functional abnormalities of this network have been found in several psychiatric conditions including autism (Kennedy et al., 2006; Monk et al., 2009). It is possible that structural abnormalities of the medial dorsal prefrontal cortex and the precuneus contribute to the memory problem and the development of psychiatric episodes in PFX+ . We also found clusters of significant negative correlations with the CGG repeat size in the dorsal medial frontal cortex (Fig. 4 and Table 4) suggesting a significant gene dosage effect of FMR1 on grey matter in this part of the brain. This further supports the influence of the CGG repeat number on the severity of clinical involvement. Past studies found particularly strong correlation of the CGG repeats with the FXTAS phenotypes including overall motor impairment (Leehey et al., 2007), the age of onset of tremor and ataxia (Tassone et al., 2007), severity of white matter disease and degree of brain atrophy (Loesch et al., 2005; Cohen et al., 2006), severity of neuropathic signs (Berry-Kravis et al., 2007b), degree of neuropathy as measured by nerve conduction studies (Soontarapornchais, 2008), reduced cerebellar volume (Adams et al., 2007), the percent of inclusions and age at death (Greco et al., 2006), and the

![Figure 5](https://brain.oxfordjournals.org/ViewPager/getImages/serials/Brain%202011%3A134%3B%20863–878%20R.-i.%20Hashimoto%20et%20al.%20Figure%205.png)
amplitude of electrophysiological response to word processing (Olichney et al., 2010).

Significant grey matter loss was also observed in the bilateral orbitofrontal cortex (Fig. 1 and Table 2). Past neuropsychological studies have shown that patients with cortical damage to this area display impulsive behaviours (Bechara et al., 2000). A recent functional MRI study of patients with obsessive–compulsive disorder and their unaffected close relatives proposed that abnormality of the orbitofrontal cortex may be endophenotype of obsessive–compulsive disorder (Chamberlain et al., 2008). Further, a recent review proposed that this area is important for subjective pleas-antness/hedonic processing (Kringlebach, 2005). Given these pro-

positions, it is tempting to associate grey matter reduction of this structure with psychiatric symptoms of disinhibition, obsessive–compulsiveness and impassivity/apathy of PFX+ (Hessl et al., 2005; Bacalman et al., 2006; Bourgeois et al., 2009). However, we did not find significant correlation of the grey matter volume and the level of obsessive–compulsiveness. It remains to be tested whether there is any association with the severity of disinhibition and apathy. We note that, in the voxel-based morphometry analysis, this area is known to be particularly sensitive to misregistra-
tion and normalization error. Therefore, we suggest that the structural abnormalities in the orbitofrontal cortex need to be replicated using different methodologies such as manual tracing (Nakamura et al., 2008) and cortical thickness measurements (Kuperberg et al., 2003).

Grey matter loss in PFX+ was clearly observed in the bilateral insula. A classic review described multiple functional roles of insu-

lar cortex including the sensory, motor and cognitive domains (Augustine, 1996). In our analysis, foci of grey matter loss were located in the posterior part, although signs of the milder grey matter loss in the anterior part were also observed when a more liberal statistical threshold was used (false discovery rate corrected \(P < 0.05\), not reported). Recent functional anatomical studies of both primates and humans highlighted the insula’s role in intero-

ception, the sense of physiological condition of the body including the visceral, hunger, pain and thermal sensations (Craig, 2003a, b). It has been proposed that altered interoception and anxiety are linked and that individuals prone to anxiety are asso-

ciated with exaggerated interoceptive prediction signals generated in the anterior insula (Paulus and Stein, 2006). Because the pos-
terior insula provides direct inputs to its anterior part, structural abnormalities of the posterior insula may adversely affect the func-
tion of the anterior insula, which contributes to anxiety-related symptoms of PFX++ and PFX−. In the present correlation analysis, although the correlation between level of obsessive–compulsiveness and reduced insula volume was significant bilaterally at an uncorrected threshold, it did not reach the significant level after correction. More observations are needed to test the possible association between the grey matter loss in the insula and anxiety-related symptoms of the premutation carriers.

Significant grey matter loss of PFX+ was also identified in the medial temporal lobe structures including parts of the fusiform gyrus and parahippocampus. Abnormalities were more extended in the left hemisphere, involving the hippocampus and amygdala. Although a number of studies indicated the pathological processes of depression in the hippocampus (Soares and Mann, 1997; Sheline, 2000), correlation between left hippocampal grey matter density and the severity of depressive symptom was marginally significant only when an uncorrected threshold was used. Previous neuropsychological and functional imaging studies have shown the critical role of the left hippocampus in verbal memory (Milner, 1972; Strange et al., 1999). Structural abnormalities of this area may therefore underlie impaired declarative verbal memory in PFX+ (Grigsby et al., 2008), although this possibility needs to be examined more directly by analysis using behavioural measures of declarative memory. In contrast, we observed a clear negative correlation between the left amygdala volume and ele-

vated levels of obsessive–compulsiveness and depression. Together with our previous functional MRI study that found functional abnormalities in the amygdala in PFX− (Hessl et al., 2007), this finding further supports the view that abnormalities in the amygdala play crucial roles in psychological symptoms of FMR1 premutation carriers. It is interesting that the left amygdala underlies both depression and an anxiety-related symptom, given a high comorbidty between depression and anxiety disorders (Kessler et al., 2003). Our result is consistent with previous findings of pathological processes affecting structure and function of the amygdala in patients with the obsessive–compulsive disorder (Szepko et al., 1999; Menzies et al., 2008) and in those with depression (Sheline, 2000).

As predicted from neuropsychological observations, significant clusters of grey matter loss of the patient group were found in multiple areas for executive cognitive functions and working memory, including the inferior frontal cortex, dorsolateral prefrontal cortex and superior parietal cortex (Figs 1 and 3; Table 2). Significant correlations between left inferior frontal cortex volume and working memory scores indicates that grey matter loss in this region, together with that in the anterior cir-
gulate cortex, contributes to working memory deficits in PFX+ and in PFX− (Grigsby et al., 2008; Cornish et al., 2009) (Fig. 3). A previous behavioural study demonstrated significant impairment in response inhibition in both affected and unaffected premutation carriers (Cornish et al., 2008). Although past brain lesion and functional imaging studies replicated the pivotal roles of the right inferior frontal cortex in response inhibition (Konishi et al., 1999; Aron et al., 2004), we did not observe significant voxels with grey matter reduction of PFX+ in this region. It still remains possible that abnormalities of this region may be more apparent in the functional measures rather than the structural ones. Furthermore, response inhibition is subserved by connectivity be-

tween the right inferior frontal cortex and other brain regions including the striatum (Aron and Poldrack, 2006). It would be interesting to examine in the future studies the structural and functional connectivity between the right inferior frontal cortex and other regions.

Although extensive grey matter atrophy was identified in PFX+, neuropsychological studies have widely reported cognitive functions that remain preserved or only mildly impaired (Cornish et al., 2008; Grigsby et al., 2008). In particular, language impairments have not been reported for any of the 16 subscales for ‘language’ and ‘verbal reasoning and comprehension’ (Grigsby et al., 2008). In the voxel-based analysis, however, a rather sig-

nificant increase of grey matter density for the patient group was
identified in the bilateral posterior part of superior/middle temporal gyrus. Although this observation has several possible explanations, the structural increase may play compensatory roles for maintaining language functions despite neurodegenerative processes affecting other language-related regions including the left inferior frontal cortex (Figs 1 and 3). Visual-spatial functions and visual attention were suggested to be only mildly impaired in PFX+ (Grigsby et al., 2008). The present voxel-based analysis revealed significant grey matter reduction in some regions of the visuospatial and dorsal visual systems, such as the right superior parietal cortex, whereas the other regions of these systems, including the right superior occipital gyrus, did not have voxels with significant grey matter reduction. It is possible that there is a functional change in intact regions that compensates for structural degeneration affecting other parts of the systems.

In addition to atrophy affecting grey matter, importance of pathological changes in the white matter has been increasingly recognized in many neurological and psychiatric conditions (Kanaan et al., 2005; Stebbins and Murphy, 2009). Indeed, hyperintensities in the periventricular and cerebellar zones have been recognized as one of the hallmark neuroradiological features of the FXTAS brain (Brunberg et al., 2002). Because the present analyses focused on alternations of grey matter, it remains unanswered how the white matter abnormalities are related to the FXTAS symptomatology and FMR1 molecular variables. The use of the MRI sequences optimal for the white matter pathology, such as the diffusion tensor imaging and the fluid-attenuated inversion recovery sequence, would complement our present findings of the grey matter abnormalities for better understanding of brain abnormalities in PFX+.

To conclude, the current voxel-based morphometry study revealed the pattern of regional grey matter abnormalities of PFX+ over the whole brain that may underlie its behavioural problems in the motor, cognitive and psychiatric domains. Findings of group comparisons together with regression analyses using FMR1 genetic molecular variables and a clinical severity scale provide foundations at the systems-level for bridging the gap between genetic molecular pathological processes and clinical behavioural observations.

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References


