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# MRI Texture Analysis Reveals Bulbar Abnormalities in Friedreich Ataxia

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## ABSTRACT

**BACKGROUND AND PURPOSE:** Texture analysis is an image processing technique that can be used to extract parameters able to describe meaningful features of an image or ROI. Texture analysis based on the gray level co-occurrence matrix gives a second-order statistical description of the image or ROI. In this work, the co-occurrence matrix texture approach was used to extract information from brain MR images of patients with Friedreich ataxia and a control group, to see whether texture parameters were different between these groups. A longitudinal analysis was also performed.

**MATERIALS AND METHODS:** Twenty patients and 21 healthy controls participated in the study. Both groups had 2 sets of T1-weighted MR images obtained 1 year apart for every subject. ROIs chosen for analysis were the medulla oblongata and pons. Texture parameters were obtained for these ROIs for every subject, for the 2 sets of images. These parameters were compared longitudinally within groups and transversally between groups.

**RESULTS:** The comparison between patients and the control group showed a significant differences for the medulla oblongata (*t* test,  $P < .05$ , Bonferroni-corrected) but did not show a statistically significant difference for the pons. Longitudinal comparison of images obtained 1 year apart did not show differences for either patients or for controls, in any of the analyzed structures.

**CONCLUSIONS:** Gray level co-occurrence matrix–based texture analysis showed statistically significant differences for the medulla oblongata of patients with Friedreich ataxia compared with controls. These results highlight the medulla as an important site of damage in Friedreich ataxia.

**ABBREVIATIONS:** C1 = the first control acquisition; C2 = the second control acquisition; FA = Friedreich ataxia; GLCM = gray level co-occurrence matrix; P1 = the first patient acquisition; P2 = the second patient acquisition

Digital images have revolutionized medicine because in addition to the readily available visual information provided by their analog counterparts, they can be mathematically manipulated by myriad processing techniques that allow extraction of many other types of information. In particular, the human eye can

only distinguish around 60 shades of gray for a given adjustment of the pupil, whereas digital medical images such as those provided by MR imaging or x-ray CT normally use around 4000 gray levels for data encoding. Texture analysis<sup>1</sup> is the name given to a set of image-processing approaches that may be used for an efficient image classification based on a reduced parameter set or to detect subtle alterations in the gray level distribution of an image or ROI. Texture analysis approaches have been widely applied in medicine to differentiate normal and pathologic tissue, such as in epilepsy,<sup>2-4</sup> Machado-Joseph disease,<sup>5</sup> and central nervous system tumors.<sup>6-8</sup> In particular, texture analysis based on the gray level co-occurrence matrix (GLCM) gives a second-order statistical description of the image or ROI.<sup>9,10</sup> In this work, the GLCM approach was used to extract information from brain MR images of patients with Friedreich ataxia (FA) and a control group. We were primarily interested in looking for differences between these groups, but we also investigated whether texture parameters correlated with clinical data in the FA group.

FA is the most common autosomal recessive ataxia, and it is

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characterized by early onset and slowly progressive gait ataxia, dysmetria, dysarthria, and deep sensory abnormalities.<sup>11</sup> It is caused by homozygous triplet *GAA* expansions in the first intron of the *FXN* gene on chromosome 9q13 in 96% of patients.<sup>12,13</sup> This leads to dramatic underexpression of the encoded protein frataxin, which is essential for proper neuronal mitochondrial functioning.<sup>11</sup> The net results of the mutation are marked and progressive neuronal loss in the dorsal root ganglia, pyramidal tracts, and dentate nuclei of the cerebellum.<sup>14</sup>

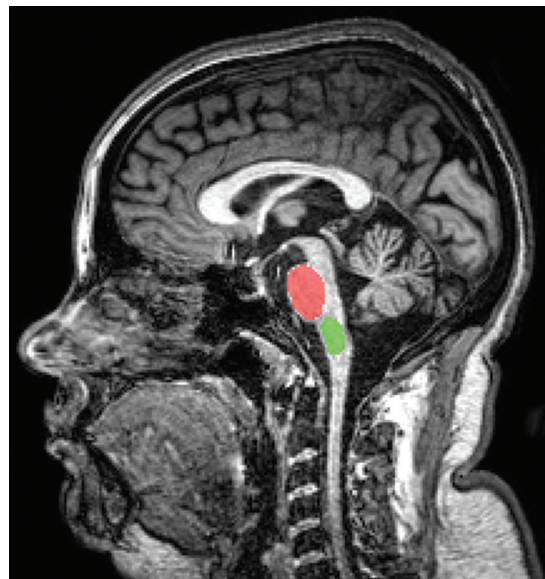
Brain MR imaging of patients with FA does not present clearly visible changes. However, some previous studies have already shown alterations by using advanced techniques. Indeed, previous voxel-based morphometry studies showed gray matter and white matter atrophy in portions of the cerebellum and brain stem in FA.<sup>15-18</sup> The spinal cord area was found to be reduced in these patients.<sup>19</sup> Also, patients with FA have WM damage in the brain stem, cerebellum, and cerebellar peduncles.<sup>16,17,20-22</sup> To the best of our knowledge, there is no previous study on GLCM texture analysis of MR imaging in FA. We believe such a study would be important because texture analysis is a powerful tool able to identify subtle abnormalities before true atrophy appears.<sup>2,3</sup> It has proved useful in finding damaged areas in closely related conditions, such as Machado-Joseph disease.<sup>5</sup> In addition, it is highly reproducible and provides quantitative results. Taken together, these characteristics make texture analysis a technique appropriate for longitudinal MR imaging-based studies, which are extremely rare in FA. In this scenario, we recruited a representative cohort of patients who underwent 2 MR imaging scans 1 year apart and performed transversal and longitudinal texture analyses. We focused the analyses on brain stem structures, the pons and medulla oblongata, because these play a key role in the pathogenesis of the disease.

## MATERIALS AND METHODS

Twenty patients with FA (mean age,  $27.6 \pm 9.6$  years; 14 women) and a control group of 21 healthy subjects (mean age,  $28.6 \pm 5.3$  years; 12 women) participated in this study. All patients had molecular confirmation of FA and were followed at the Neurogenetics Clinic at the University of Campinas. Mean ages at diagnosis and disease duration were  $15.4 \pm 5.2$  and  $13.1 \pm 8.4$  years, respectively. We used the Friedreich ataxia rating scale at each visit to quantify disease severity for each patient.<sup>23</sup> The study was approved by the local ethics committee, and all participants signed an informed consent form, before data acquisition.

MR imaging data were acquired in a 3T Achieva scanner (Philips Healthcare, Best, the Netherlands) at the Clinics Hospital of our university. Images used for texture analysis were T1-weighted, acquired with an 8-channel head coil, with an FOV of  $240 \times 240 \times 180$  mm<sup>3</sup>, voxel size of  $1 \times 1 \times 1$  mm<sup>3</sup>, TR of 7 ms, and TE of 3.2 ms.

Given an ROI in an image and a pixel distance in a given direction (eg, 2 pixels in the vertical direction), the GLCM approach consists of calculating, for every possible pair of gray levels, how many times they co-occur in the ROI in this direction and distance. From a GLCM, statistical parameters can then be computed, the most usual ones being the following<sup>24</sup>: contrast, homogeneity, correlation, variance, sum average, sum variance, differ-



**FIG 1.** Example of segmented ROIs in the center section of the MR imaging examination for a patient with FA. The region in red indicates the pons, and the region in green indicates the medulla oblongata.

ence variance, uniformity, entropy, sum entropy, and difference entropy. Formulas and descriptions for these parameters can be found in the On-line Table.

As mentioned, the ROIs investigated in this work were the medulla oblongata and the pons, known to be affected by FA. These structures were manually segmented, by using MaZda software (<http://www.eletel.p.lodz.pl/programy/mazda/>),<sup>25</sup> in 7 sagittal sections for each subject (consisting of the central section plus 3 anterior and 3 posterior sections). Figure 1 shows an example of these ROIs, segmented in 1 section, for a patient with FA, where the region in red is the pons and the region in green is the medulla oblongata. GLCMs were computed for every section, also by using the MaZda software. This software computes GLCMs for 4 directions (horizontal, vertical, and 2 diagonals) and 5 distances (1–5 pixels), giving 20 GLCMs per section or, in the present case, 140 GLCMs per ROI (280 per subject). Because the aforementioned 11 parameters were computed for every GLCM, the total number of parameters was 1540 per ROI (3080 per subject).

To reduce the number of parameters, we computed 2 averages. First, an average over the 7 ROI sections was calculated for each parameter, weighted by the segmented area in each section; this procedure had the effect of “merging” the information obtained for a given ROI that was previously “sliced.” Second, an average over the 4 directions was calculated for each parameter. The idea behind this approach was to make the parameters direction-independent, because even if a given tissue had a preferred structural direction, the position axes of the acquired MR images might be slightly different for different subjects; this difference could affect the results. After these procedures, parameters were reduced to 55 per ROI (110 per subject).

Finally, GLCM texture parameters were compared statistically, by using a *t* test, Bonferroni corrected, with a significance level of  $\alpha = .05$ . Each GLCM parameter was analyzed longitudinally within a group, comparing images taken 1 year apart, and transversally between the groups. For the longitudinal compari-

**GLCM texture parameters, which were significantly different between patients with FA and controls for at least 1 comparison between groups<sup>a</sup>**

GLCM Distance (pixel)	Parameter	Group (Mean)				Comparison (P Value, Bonferroni-Corrected, t Test)			
		P1	P2	C1	C2	P1 × C1	P1 × C2	P2 × C1	P2 × C2
Medulla oblongata									
1	Correlation	0.27 ± 0.03	0.25 ± 0.05	0.45 ± 0.09	0.41 ± 0.10	.000 <sup>b</sup>	.000 <sup>b</sup>	.000 <sup>b</sup>	.000 <sup>b</sup>
1	Variance	11.84 ± 3.36	12.80 ± 4.14	17.71 ± 5.10	15.60 ± 4.83	.008 <sup>b</sup>	.188	.413	1.00
1	Sum average	86.25 ± 3.45	86.18 ± 2.62	110.75 ± 5.83	103.48 ± 14.32	.000 <sup>b</sup>	.002 <sup>b</sup>	.000 <sup>b</sup>	.002 <sup>b</sup>
1	Sum variance	32.55 ± 9.52	34.61 ± 10.86	50.24 ± 16.52	43.56 ± 14.79	.014 <sup>b</sup>	.103	.484	1.00
1	Entropy	1.46 ± 0.15	1.47 ± 0.07	2.00 ± 0.08	1.86 ± 0.23	.000 <sup>b</sup>	.000 <sup>b</sup>	.000 <sup>b</sup>	.000 <sup>b</sup>
1	Sum entropy	0.98 ± 0.08	0.99 ± 0.04	1.36 ± 0.06	1.25 ± 0.17	.000 <sup>b</sup>	.000 <sup>b</sup>	.000 <sup>b</sup>	.000 <sup>b</sup>
1	Difference entropy	0.71 ± 0.04	0.72 ± 0.04	0.93 ± 0.05	0.86 ± 0.11	.000 <sup>b</sup>	.000 <sup>b</sup>	.000 <sup>b</sup>	.002 <sup>b</sup>
1	Homogeneity	0.21 ± 0.05	0.20 ± 0.02	0.27 ± 0.03	0.27 ± 0.04	.004 <sup>b</sup>	.165	.000 <sup>b</sup>	.000 <sup>b</sup>
2	Correlation	0.05 ± 0.05	0.04 ± 0.06	0.18 ± 0.08	0.17 ± 0.10	.000 <sup>b</sup>	.001 <sup>b</sup>	.000 <sup>b</sup>	.002 <sup>b</sup>
3	Correlation	0.04 ± 0.05	0.02 ± 0.05	0.13 ± 0.07	0.14 ± 0.08	.002 <sup>b</sup>	.004 <sup>b</sup>	.000 <sup>b</sup>	.000 <sup>b</sup>
4	Correlation	-0.01 ± 0.04	-0.03 ± 0.05	0.08 ± 0.07	0.08 ± 0.08	.004 <sup>b</sup>	.013 <sup>b</sup>	.000 <sup>b</sup>	.001 <sup>b</sup>
5	Correlation	-0.02 ± 0.03	-0.03 ± 0.05	0.06 ± 0.07	0.05 ± 0.07	.006 <sup>b</sup>	.055	.008 <sup>b</sup>	.049 <sup>b</sup>
Pons									
1	Difference variance	7.27 ± 2.24	7.56 ± 1.89	5.58 ± 1.45	5.38 ± 1.11	.490	.142	.125	.034 <sup>b</sup>

<sup>a</sup> Both patient acquisitions were compared with both control acquisitions.

<sup>b</sup> Significant.

son, only 16 (of 20) patients had the corresponding MR imaging acquisitions; therefore, these analyses were made considering this reduced group (16 patients). In addition, GLCM parameters that were found significant were correlated to clinical variables (age, age at onset, disease duration, GAA1, GAA2, Friedreich Ataxia Rating Scale total score) within the patient group. A *t* test comparing the age of the groups was also performed to ensure that ages were matched.

**RESULTS**

We will refer to P1 as the group corresponding to the first set of images (20 images) of the patients with FA, P2 as the group corresponding to the second set (16 images, obtained 1 year later) of the patients with FA, and C1 and C2 correspond to the first and second sets of control acquisitions, respectively (21 images each, obtained 1 year apart).

The FA and control groups were not significantly different regarding age (*t* test; *P* = .99 for P1 × C1; *P* = .69 for P1 × C2; *P* = .60 for P2 × C1; *P* = .86 for P2 × C2).

Longitudinal comparison of GLCM texture parameters for images obtained 1 year apart did not show significant alterations for either the control group (C1 × C2, *t* test, *P* > .05 for all parameters) or the FA group (P1 × P2, *t* test, *P* > .05 for all parameters) for the analyzed structures (medulla oblongata and pons).

For the transversal study, because there were no significant age differences between groups, both patient acquisitions were compared with both control acquisitions, to verify the reproducibility of the results. Significant changes between patients with FA and controls were found for many GLCM parameters for the medulla oblongata, for all comparisons (P1 × C1, P1 × C2, P2 × C1, P2 × C2); and only 1 parameter, for 1 comparison (P2 × C2), showed significant differences for the pons. These results are summarized in the Table. In this Table, only parameters that were significant for at least 1 comparison are shown. All remaining parameters were not significant, for any comparison.

There were no significant correlations between significant GLCM parameters and clinical variables.

**DISCUSSION**

The aim of the present work was to investigate brain stem damage in FA by using texture analysis of T1-weighted images. Indeed, texture analysis based in the GLCM is a technique capable of extracting a set of parameters that characterize the gray level distribution of the analyzed ROIs. In turn, the gray level distribution reflects physical properties of the underlying tissues, in this case related to the local proton attenuation and T1 constant of the tissue, because T1-weighted MR images were used. In addition, because the resolution of the images used was 1 mm<sup>3</sup>, the reflected tissue properties are an average of microscopic characteristics (see De Certaines et al<sup>26</sup> for a discussion on the limitations of correlating texture analysis results to histology). Nevertheless, obtained results showed significant alterations for patients with FA compared with controls, mainly for the medulla oblongata. In particular, the results for the medulla were very robust because most of the significant parameters were robust for all the comparisons made between the different sets of images obtained from patients with FA and controls. Conversely, the result found for the pons, in which only 1 parameter was significant (difference variance) for only 1 of the comparisons (P2 × C2), may be regarded as a highly probable spurious finding. Most interesting, there were no alterations found in the longitudinal study. This could be because the time span (1 year) was too small for noticeable changes in such a slowly evolving disorder. In fact, in analyses of a large set of control images, no changes with age were found in texture parameters for the medulla oblongata or pons (unpublished data, T.A.S. and G.C., 2014).

We believe that the texture abnormalities found here reflect bulbar structural changes, but not exactly atrophy of the structure. This hypothesis is supported by a previous texture-based study in patients with a related neurodegenerative disorder, namely amyotrophic lateral sclerosis, that identified prominent

texture abnormalities earlier than true volumetric reduction.<sup>27</sup> Also in line with these findings, some previous volumetric studies in FA failed to identify bulbar atrophy,<sup>17</sup> and others actually found it,<sup>15</sup> but it was restricted to the dorsal part of the structure in such a way that it would be difficult to explain our consistent texture modifications. Therefore, we opted not to include volumetric measurements of the medulla as a covariate in our statistical approach to compare groups.

The GLCM parameters that were significant for the medulla oblongata can be roughly divided into 3 groups, according to “the purpose of the weights in the corresponding equations.”<sup>28</sup> These groups can help chart a slightly more intuitive meaning of the findings.

One group is related to descriptive statistical measures of the GLCM<sup>28</sup> and includes the parameters correlation, variance, sum average, and sum variance. In particular, the correlation parameter, which gives a measure of the linear dependency between the gray levels of co-occurring (separated by the given distance) pixels, was consistently significant for the medulla oblongata for all GLCM distances for all comparisons, with the only exception being distance 5 for P1 × C2, though in this case, the *P* value was very close to significance (*P* = .055, Table). It has already been shown that this measure has the same expectation as the autocorrelation coefficient, particularly when the distance between pixels used in the GLCM calculation is small compared with the ROI dimensions,<sup>29</sup> which was the case here for the smallest GLCM distances (1 and 2 pixels) (Appendix). The correlation was smaller for patients compared with controls, for all distances; this difference therefore indicates a decrease in the autocorrelation coefficient for the gray level distribution of the medulla oblongata for the patients with FA. In other words, the gray level distribution of this ROI is less stationary or less periodic for patients than for controls.

The variance parameter measures the dispersion of the gray level distribution; the sum average and sum variance parameters measure, respectively, the mean and the dispersion of the distribution related to the sum of the co-occurring gray levels (On-line Table). From these 3 parameters, only the sum average was significant for all comparisons (Table), with variance and sum variance only significant for the comparison P1 × C1. All these parameters were smaller for patients with FA compared with controls. This finding therefore indicates a smaller mean and perhaps a tendency to a decrease in dispersion of the co-occurring gray level values for the medulla oblongata of patients with FA.

In another group are the entropy measures (entropy, sum entropy, and difference entropy). Entropy is a widely known measure coming from information theory,<sup>30</sup> which can be used to evaluate the amount of orderliness or the information content of a dataset. The greater the entropy, the more random is the underlying distribution (in this case, the distribution of the co-occurring gray levels or the distribution of the sum of the co-occurring gray levels or the distribution of the difference of the co-occurring gray levels). In the present analysis, all these measures were decreased for patients with FA compared with controls. In terms of the gray level distributions (both the original between co-occurring pixels and the sum and difference distributions), the decrease

of the entropy measures points to the appearance of some sort of orderliness in the data of patients with FA for the medulla oblongata, which was not there in the control data.

Finally, in the last group, related to contrast measures, which “use weights related to the distance from the GLCM diagonal,”<sup>28</sup> is homogeneity, which, as shown by its name, is related to the homogeneity of the ROI. This parameter was significant for 3 of the comparisons (the only exception being P1 × C2, Table), and it decreased for patients with FA compared with controls.

Together, these results indicate probable changes in the underlying tissue composing the medulla oblongata. Although, to the best of our knowledge, there are no studies correlating in vivo texture measures with histopathology, Zhang et al<sup>31</sup> performed a study in which they found correlation among histopathology and texture parameters obtained from MR imaging of postmortem tissue samples of patients with multiple sclerosis. Possible tissue changes in the medulla oblongata are in line with previous pathologic reports in FA. In fact, medullar atrophy has been described since the original descriptions of Nikolaus Friedreich.<sup>32</sup> This atrophy is mostly due to neuronal loss and shrinkage of gracile and cuneate nuclei, which are related to the dorsal sensory pathways in the spinal cord. In addition, corticospinal tracts are also damaged at this point.<sup>14</sup> The lack of correlation between texture parameters and clinical data was somewhat unexpected but might be related to our segmentation strategy. We opted to use large ROIs that included the whole medulla oblongata, and neurodegeneration in FA is not homogeneous in this region (posterior structures are more severely damaged). Future studies using smaller ROIs within the medulla might prove useful in this setting.

## CONCLUSIONS

In the present work, we found statistically significant differences for the medulla oblongata of patients with Friedreich ataxia compared with healthy individuals, by using GLCM-based texture analysis. These results highlight the medulla as an important site of damage in FA.

## APPENDIX

ROI dimensions decrease with GLCM distance. In this work, the mean ROI areas used ranged approximately from 110 pixels (for a distance of 5 pixels) to 240 pixels (for a distance of 1 pixel) for the medulla of the patients with FA; from 160 to 300 pixels for the medulla of the controls; from 380 to 570 pixels for the pons of the patients with FA; and from 320 to 500 pixels for the pons of the controls. The area distributions were not significantly different between groups for any of the ROIs.

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