

Spatial patterns of brain atrophy in MCI patients, identified via high-dimensional pattern classification, predict subsequent cognitive decline

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Spatial patterns of brain atrophy in mild cognitive impairment (MCI) and Alzheimer's disease (AD) were measured via methods of computational neuroanatomy. These patterns were spatially complex and involved many brain regions. In addition to the hippocampus and the medial temporal lobe gray matter, a number of other regions displayed significant atrophy, including orbitofrontal and medial-prefrontal grey matter, cingulate (mainly posterior), insula, uncus, and temporal lobe white matter. Approximately 2/3 of the MCI group presented patterns of atrophy that overlapped with AD, whereas the remaining 1/3 overlapped with cognitively normal individuals, thereby indicating that some, but not all, MCI patients have significant and extensive brain atrophy in this cohort of MCI patients. Importantly, the group with AD-like patterns presented much higher rate of MMSE decline in follow-up visits; conversely, pattern classification provided relatively high classification accuracy (87%) of the individuals that presented relatively higher MMSE decline within a year from baseline. High-dimensional pattern classification, a nonlinear multivariate analysis, provided measures of structural abnormality that can potentially be useful for individual patient classification, as well as for predicting progression and examining multivariate relationships in group analyses.

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Introduction

Alzheimer's disease (AD) is the most common dementia, with incidence rates doubling every 5 years after the age of 65. It is estimated that half of the population above 80 years may have symptomatic AD, and that this number will grow rapidly as life expectancy increases, and as the baby boomers' generation moves into the high risk age group. The psychological and financial cost of AD is tremendous and rapidly rising. Although there is currently no disease-modifying treatment, many potential treatments are being tested, some of which may have significant side-effects. It is therefore becoming clear that effective and well-targeted treatment necessitates early diagnosis of the disease.

Currently, definitive diagnosis of AD can be made if an autopsy documents the presence of the characteristic neuritic β -amyloid plaques and neurofibrillary tangles in the appropriate brain regions in an individual with a history of progressive dementia. Therefore, there has been a keen interest in the neuroimaging community to develop imaging-based biomarkers, especially of early AD stages (Braak et al., 1998), as well as for predicting individuals that are likely to progress to AD and are therefore good candidates for therapy. Magnetic resonance imaging (MRI) can potentially play an important role as diagnostic tool, mainly because it is widely available and part of the American Academy of Neurology standard clinical evaluation for individuals with symptoms of dementia. MRI helps measure spatial patterns of atrophy, and their evolution with disease progressions, which are surrogate markers of the underlying neurodegenerative AD pathology.

The neuroimaging literature is rich in studies measuring volumes of regions of interest (ROIs) known to be affected by AD, especially of the hippocampus and the entorhinal cortex (Kaye et al., 1997; Jack et al., 1999; Convit et al., 2000; Killiany et al., 2000; Dickerson et al., 2001; Chetelat et al., 2002; Visser et al., 2002; Stoub et al., 2005; De Leon et al., 2006); more complex

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Available online on ScienceDirect (www.sciencedirect.com).

shape properties of hippocampal ROIs have also been investigated (Csernansky et al., 2005). However, the pattern of AD pathology is complex and evolves as the disease progresses, starting mainly in the hippocampus and entorhinal cortex, and subsequently spreading throughout most of the temporal lobe and the posterior cingulate, finally involving extensive cortical regions, especially parietal, prefrontal, and orbitofrontal. Therefore, measuring volumes of a few structures cannot capture the spatio-temporal pattern of brain atrophy in its entirety. Moreover, measurements of hand-drawn ROIs are not easily reproducible within and across different raters. Finally, the pattern of atrophy associated with AD does not necessarily follow pre-determined anatomical boundaries.

During the past decade, methods of computational neuroanatomy, such as voxel-based and deformation-based analysis, have gained attention in the neuroimaging community (Davatzikos et al., 2001; Thompson et al., 2001; Chetelat et al., 2002; Ashburner et al., 2003; Karas et al., 2004; Pennanen et al., 2005; Bozzali et al., 2006; Saykin et al., 2006; Xie et al., 2006; Whitwell et al., 2007), because they allow for the complete evaluation of structural and functional brain images, without the need to make *a priori* assumptions about the size, extent, and number of regions to be measured. Instead, these methods apply voxel-by-voxel evaluation of the images, and identify potentially complex spatial patterns of brain atrophy.

In addition to voxel-based analysis methods, techniques for high-dimensional pattern classification have begun to find their way to the literature of neuroimaging of AD (Lao et al., 2003; Lao et al., 2004; Liu et al., 2004; Adeli et al., 2005; Tandon et al., 2006; Li et al., 2007), aiming to provide computational tools that classify individuals, based on their MRI or PET scans, rather than determining statistical group differences. The current study builds upon previous work in Davatzikos et al. (in press), which used a limited sample of patients with MCI and cognitively healthy individuals to construct classifiers that separate the two groups. The current study, however, emphasizes (1) application to a larger sample from the ADNI study; (2) inclusion of AD patients, in addition to healthy and MCI individuals; (3) a different methodological design, in which structural differences between healthy individuals and AD patients are used to construct a high-dimensional classifier, which is subsequently applied to MCI patients, rather than emphasizing differences between MCI and controls. This approach allows us to determine MCI subgroups that have structural profiles similar to AD or to healthy individuals. Most importantly, it allowed us to further associate these structural profiles with Mini Mental State Examination (MMSE) scores and their 1-year change in follow-up examinations, and demonstrate their prognostic value, an issue of very high importance currently in the AD literature.

In particular, the current study pursues a voxel-based morphometric analysis of cognitively normal individuals, individuals with MCI, and AD patients, using an atlas warping approach used to generate regional tissue density maps that reflect the regional distribution of brain tissue. The hypothesis was that this approach would allow us to quantitatively capture complex spatial pattern of brain atrophy that can potentially serve as sensitive and specific imaging signatures of MCI and AD. The classification analysis also offers one possible way to classify an entire pattern of atrophy to AD or cognitively normal individuals (CN), and potentially to predict whether an MCI subject will eventually develop AD, using longitudinal follow-ups.

Materials and methods

Data used in the preparation of this article were obtained from the Alzheimer's Disease Neuroimaging Initiative (ADNI) database (www.loni.ucla.edu/ADNI). The goal of ADNI is to recruit 800 adults, ages 55 to 90, to participate in the research—approximately 200 CN older individuals to be followed for 3 years, 400 people with MCI to be followed for 3 years, and 200 people with early AD to be followed for 2 years. For up-to-date information, see www.adni-info.org.

Participants

All ADNI participants with structural MR images available on the ADNI Web site as of February 2007 (the latest scan was from December 19, 2006) were part of this analysis. This included 66 CN individuals (mean age \pm S.D., 75.18 \pm 5.39), 88 MCI patients (76.38 \pm 7.60), and 56 AD patients (77.40 \pm 7.02), whose MRI scans were analyzed. The MMSE scores (mean \pm S.D.) of each group at baseline were 29.08 \pm 0.97, 26.78 \pm 1.91, and 23.07 \pm 1.83, respectively. The groups were relatively well-balanced in terms of gender (50%, 36%, 57% women in each of the 3 groups, respectively). MMSE scores from the subset of participants that had completed 3 follow-up exams by the end of this study in June 2007 were also included in the analysis, and used as a measure of disease progression.

Images

The data sets included standard T1-weighted MR images acquired sagittally using volumetric 3D MPRAGE with 1.25 \times 1.25 mm in-plane spatial resolution and 1.2 mm thick sagittal slices (8° flip angle). Most of the images were obtained using 1.5 T scanners, while a few were obtained using 3T scanners: 8 CN, 11MCI, and 8 AD patients. Detailed information about MR acquisition procedures is available at the ADNI Web site.

Image analysis

Images were first preprocessed according to previously validated and published techniques (Goldszal et al., 1998). The pre-processing steps included (1) alignment to the AC-PC plane; (2) removal of extra-cranial material (skull-stripping); (3) tissue segmentation into grey matter (GM), white matter (WM), and cerebrospinal fluid (CSF), using a brain tissue segmentation method proposed in Pham and Prince (1999); (4) high-dimensional image warping (Shen and Davatzikos, 2002) to a standardized coordinate system, a brain atlas (template) that was aligned with the MNI coordinate space (Kabani et al., 1998); (5) formation of regional volumetric maps, named RAVENS maps (Goldszal et al., 1998; Davatzikos et al., 2001; Shen and Davatzikos, 2003), using tissue preserving image warping (Goldszal et al., 1998). RAVENS maps quantify the regional distribution of GM, WM, and CSF, since one RAVENS map is formed for each tissue type. In particular, if the image warping transformation that registers an individual scan with the template applies an expansion to a GM structure, the GM density of the structure decreases accordingly to insure that the total amount of GM is preserved. Conversely, a RAVENS value increases during contraction, if tissue from a relatively larger region is compressed to fit a smaller region in the template. Consequently, RAVENS values in the template's (stereotaxic) space are directly proportional to the volume of the

respective structures in the original brain scan. Therefore, regional volumetric measurements and comparisons are performed via measurements and comparisons of the respective RAVENS maps. For example, patterns of GM atrophy in the temporal lobe are quantified by patterns of RAVENS decrease in the temporal lobe in the stereotaxic space.

The RAVENS approach has been extensively validated (Goldszal et al., 1998; Davatzikos et al., 2001) and applied to a variety of studies (Resnick et al., 2000, 2001; Kim et al., 2003; Resnick et al., 2003, 2004; Beresford et al., 2006a,b; Gur et al., 2006; Stewart et al., 2006; Driscoll et al., 2007). It bears similarities with the “optimized VBM” approach (Good et al., 2002), except it uses a highly conforming high-dimensional image warping algorithm that captures finer structural details. Moreover, it uses tissue-preserving transformations, which ensures that image warping absolutely preserves the amount of GM, WM, and CSF tissue present in an individual’s scan, thereby allowing for local volumetric analysis.

Statistical analysis and pattern classification

Group comparisons were performed via voxel-based statistical analysis of respective RAVENS maps that were normalized by intra-cranial volume and smoothed using 8 mm full-width at half-maximum (FWHM) smoothing kernel. Group comparisons involved voxel-by-voxel *t*-tests applied by the SPM2 software (<http://www.fil.ion.ucl.ac.uk/spm/software/spm2>). Comparison for multiple corrections utilized the false discovery rate (FDR) method (Yekutieli and Benjamini, 1999), as implemented in the SPM software. In addition to the group analyses, we perform individual-patient analysis, aiming to classify individual scans belonging to CN, MCI, or AD participants. This analysis is important because it directly relates to our ability to use quantitative MRI analysis for individual diagnosis, rather than to identify statistical differences between two potentially overlapping groups. Toward this end, we applied a high-dimensional pattern classification approach, which has been published and used in various neuroimaging studies (Fan et al., 2005, 2007, in press; Davatzikos et al., in press). This approach considers all brain regions jointly, and identifies a minimal set of regions whose volumes jointly maximally differentiate between the two groups under consideration, on an individual scan basis. Leave-one-out cross-validation is used to test this classification scheme on data sets not used for training, and obtain a relatively unbiased estimate of the generalization power of the classifier to new patients. The pattern classification method provides a *structural phenotypic score* (SPS). For a classifier constructed from the CN and AD groups, positive SPS implies AD-like brain structure, and vice versa. The classifier that was determined to maximally distinguish between CN and AD participants was subsequently applied to the MCI group.

We also utilized the standard region of interest (ROI) method to analyze the volumes of the hippocampus and the entorhinal cortex via a template warping method that has been previously published and validated (Shen and Davatzikos, 2002; Shen et al., 2002), in order to determine whether conventional ROI measurements are sufficient for classification of individual scans with high sensitivity and specificity.

SPS and rates of MMSE change

Since at this stage of the study, the clinical outcome is not yet available for most of the participants, we evaluated associations

between the SPS determined from the MRI and rates of changes of the MMSE scores, which were calculated for those MCI individuals ($n=38$) that had at least 3 examinations (baseline plus 2 follow-ups, 12 months). These rates of change were computed separately for the two subgroups identified by the pattern classification: the MCI_CN (MCIs having CN-like patterns, $n=16$) and MCI_AD (MCIs having AD-like patterns, $n=22$) subgroups. Conversely, we divided the MCI participants into two groups, the progressors (MCI_PR) and the nonprogressors (MCI_NPR). (We stress that strictly speaking, this categorization does not reflect progression to AD, but progression of the MMSE scores toward relatively lower values.) We then evaluated whether the pattern classification approach can separate MCI_PR from MCI_NPR, knowing that this is an extremely difficult task due to the noise that is inherent to the measurement of rate of change of MMSE, especially from 3 measurements within 1 year, but also because short-term MMSE decline does not necessarily imply clinical progression to AD. One of the caveats in this analysis was that it is difficult to define a threshold for MMSE rate of change that would define the subgroups MCI_PR and MCI_NPR. We cannot use a threshold of 0, because we know that even CN individuals display some decline. Therefore, we decided to examine a range of possible thresholds on MMSE rate of change, and test the group separability within that range. The hypothesis was that very low or very high thresholds would lead to nonseparable subgroups (since they would lump together progressors and nonprogressors into the same class), and somewhere in the small negative range (small rate of MMSE change) we should find maximal separability.

Results

Region of interest volumetry

The volumes of the hippocampus (left+right) against the entorhinal cortex (left+right), after normalization by intra-cranial volume (ICV), are shown in Fig. 1 as a scatter plot. AD and CN are relatively well separated, although classification accuracy would not be clinically sufficient, in terms of providing adequate sensitivity and specificity on an individual patient basis. Volumes of the MCI group completely overlap with both groups, especially the AD group. We also constructed SVM classifiers using these volume measurements. The classification accuracy computed via the leave-one-out cross-validation was 82.0%, 76.0%, and 58.3%, for AD vs. CN, MCI vs. CN, and AD vs. MCI, respectively.

Voxel-based analysis of RAVENS maps

Statistically significant results from the CN vs. AD comparisons via voxel-based statistical analysis are shown in Fig. 2. These maps display the *t*-statistic of voxel-wise *t*-tests; however, only the clusters with *p* values corrected for multiple comparison above $p=0.05$ were obtained. SPM2 (FDR multiple comparison correction) was used for all voxel-based analyses. Fig. 2 indicates severe GM atrophy in the AD group, and less pronounced WM atrophy mainly located around the hippocampus. Apparent WM atrophy in the anterior periventricular region is due to periventricular leuko-areosis that tends to be segmented as GM, due to its darker T1 signal.

Statistically significant findings from the voxel-based comparisons between the CN and MCI groups are shown in Fig. 3. The pattern of atrophy is similar to the one in Fig. 2, except less pronounced, as indicated by the values of the *t*-statistic. No significant

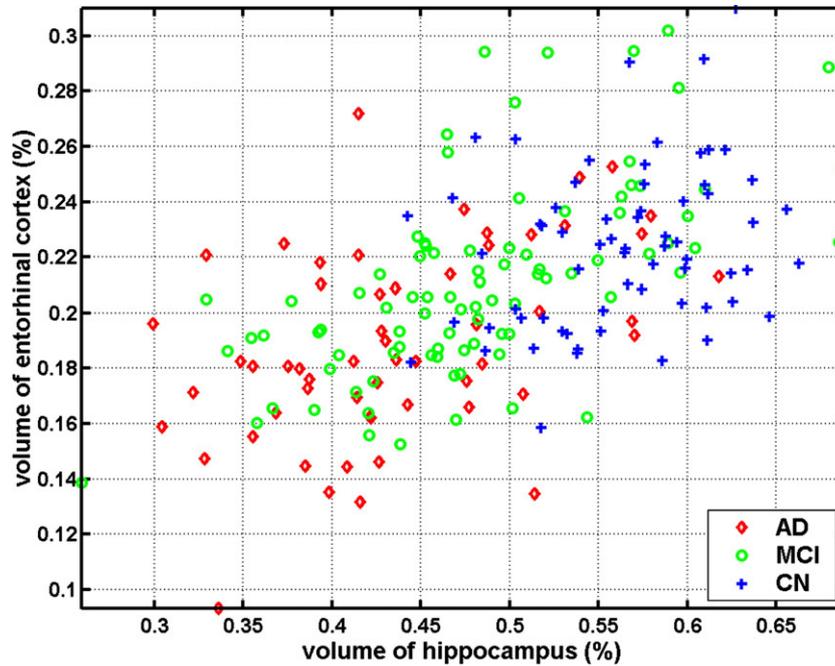


Fig. 1. Scatter plot of the volumes of the hippocampus (left+right) against the entorhinal cortex (left+right) of the groups of CN, MCI, and AD, after normalization by ICV.

WM group differences were found, after correction for multiple comparisons, therefore the WM regions shown were determined without multiple comparison correction at the $p < 0.001$ level.

Fig. 4 shows the statistically significant findings from the MCI vs. AD comparison. Even smaller regional volumetric differences were found in this group comparison. Since the differences between MCI and AD were relatively small and almost disappeared after correction for multiple comparisons, Fig. 4 also displays the results prior to FDR correction.

Pattern classification

The pattern classification approach was initially applied separately to each group comparison: (1) AD vs. CN; (2) MCI vs. CN; and (3)

AD vs. MCI. The classification accuracy was determined via the leave-one-out (LOO) cross-validation to be 94.3%, 81.8%, and 74.3%, respectively, for the 3 comparisons. Because LOO was applied, these are estimations of classification accuracy of a new individual’s scan and therefore of direct diagnosis relevance. These classifiers’ receiver operating characteristic (ROC) curves are shown in Fig. 5 (these ROC curves were determined using LOO). These classifications’ respective AUCs (area under the ROC curve) were 96.5%, 84.6%, and 75.9%.

Subgrouping of MCI participants and follow-up MMSE scores

In order to further investigate the patterns of brain atrophy in the MCI participants, the classifier built from the AD and CN groups was applied to MCI participants. As described in Materials

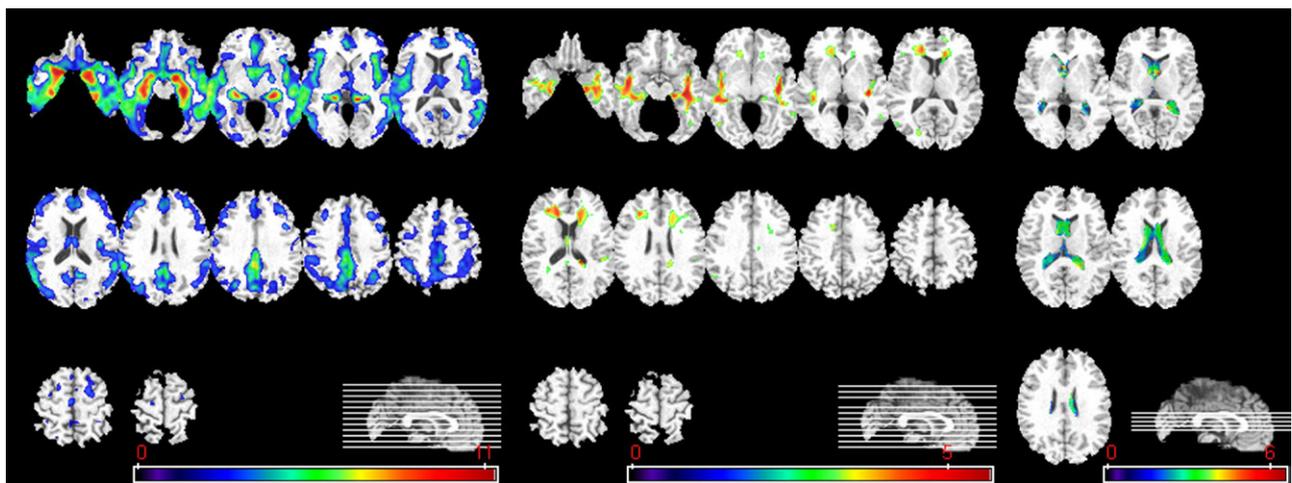


Fig. 2. Voxel-based analysis of group difference between CN and AD. From left to right, group comparison results on GM, WM, and CSF are shown (GM, WM: CN > AD, CSF: AD > CN, $p < 0.05$, corrected). The color-maps indicate the scale for the t -statistic. Images are displayed in radiological convention.

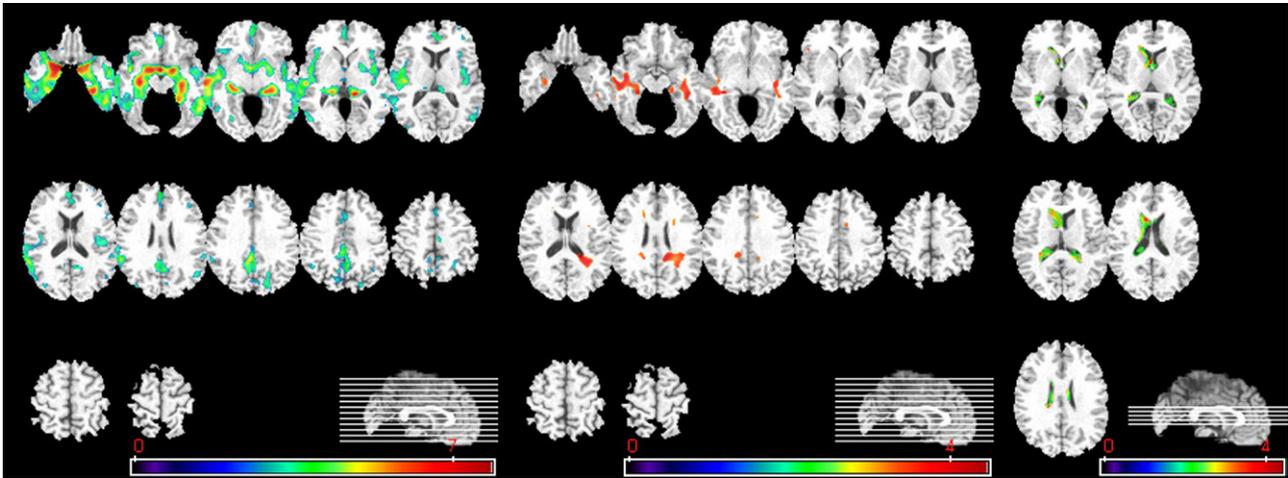


Fig. 3. Voxel-based analysis of group difference between CN and MCI. From left to right, group comparison results on GM, WM, and CSF are shown (GM: CN>MCI, $p < 0.05$, FDR-corrected; CSF: MCI>CN, $p < 0.05$, FDR-corrected; WM: CN>MCI, $p < 0.001$, uncorrected). The color-maps indicate the scale for the t -statistic. Images are displayed in radiological convention.

and methods, this classifier provides SPS that is positive for AD-like structure and negative for CN-like structure. Fig. 6 shows the distribution of the SPSs of all MCI participants, further indicating that the majority of the MCI participants displayed AD-like structural profiles. MCI participants were further divided into the ones that had positive SPS (AD-like patterns) and the ones that had negative SPS (CN-like patterns). We refer to these two groups as MCI_{AD} (57 MCI participants) and MCI_{CN} (31 MCI participants), respectively. These two subgroups were then compared via voxel-based analysis of their RAVENS maps. Statistically significant regional volumetric differences are shown in Fig. 7.

Fig. 8 shows the group differences between MCI_{AD} and CN participants via voxel-based statistical analysis. It is worth noting that the group differences in WM RAVENS maps were even stronger than group differences between AD and CN participants.

Fig. 9 shows the group differences between MCI_{CN} and AD participants via voxel-based statistical analysis. The group differences between MCI_{CN} and AD participants are similar with those between CN and MCI_{AD}. Almost no significant group differences were found between MCI_{CN} and CN participants via voxel-based statistical analysis: only a small region in the medial orbitofrontal cortex passed the $p = 0.05$ threshold, after FDR correction for multiple comparisons. Similarly, no group differences between MCI_{AD} and AD participants reached significance, after FDR correction.

SPS and rate of MMSE change

As discussed in Materials and methods, rates of change of the MMSE scores were calculated separately for the two subgroups

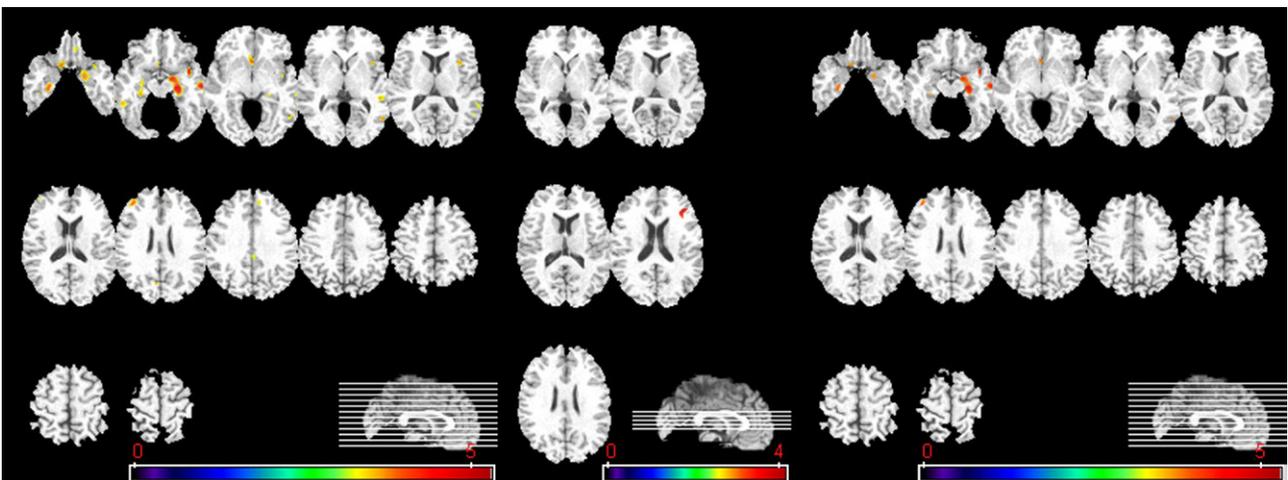


Fig. 4. Voxel-based analysis of group difference between MCI and AD. Left column shows GM comparisons, and middle column shows WM comparisons, without correction of multiple comparisons (MCI>AD, $p < 0.001$, uncorrected). After FDR correction (MCI>AD, $p < 0.05$), significant group differences are found only in GM RAVENS maps, as shown in right column. No significant group difference was found on CSF comparisons. The color-maps indicate the scale for the t -statistic. Images are displayed in radiological convention.

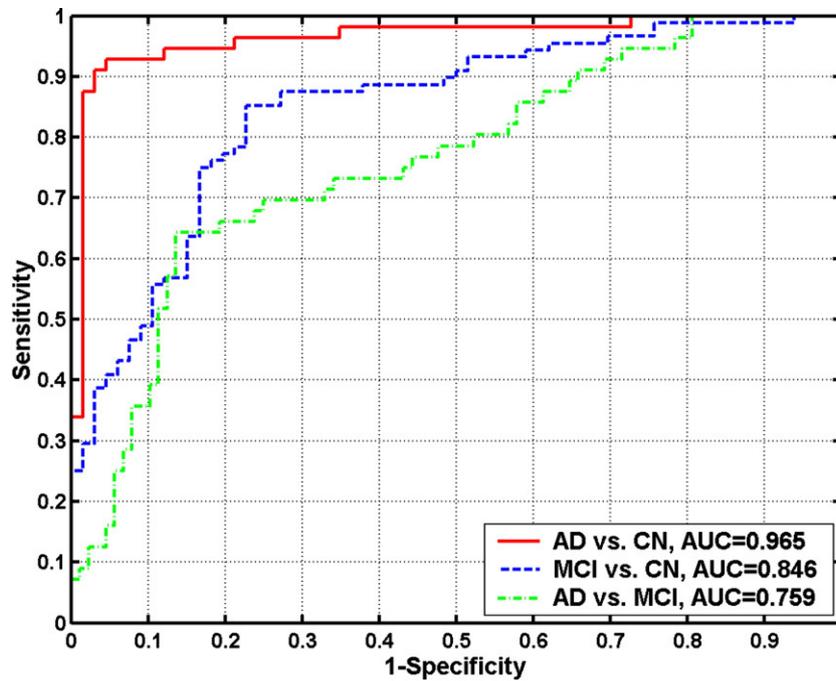


Fig. 5. ROC curves showing the overall classification performance in MRI-based classification of AD from CN, MCI from CN, and AD from MCI. Their respective AUCs (area under the ROC curve) are 0.965, 0.846, and 0.759.

identified by the pattern classification: the MCI_{CN} ($n=16$) and MCI_{AD} ($n=22$) subgroups. The rate of MMSE score annual decrease of the former group was significantly smaller than that of the latter group with a p value of 0.028. The average rate of MMSE score annual decrease (mean \pm S.D.) was -0.30 ± 3.13 for the former group and -2.31 ± 3.07 for the latter group. The correlation coefficient between the relationship of the SPS and the rate of

MMSE change was -0.39 ($p=0.0155$). The distributions of the scores during baseline and follow-ups are shown in Fig. 10, and a regression plot in Fig. 11.

In Materials and methods, we discussed the categorization of the MCI cohort into progressors and nonprogressors: MCI_{PR} and MCI_{NPR}, via thresholding their rates of MMSE change at various thresholds. The classification results, after leave-one-out cross-

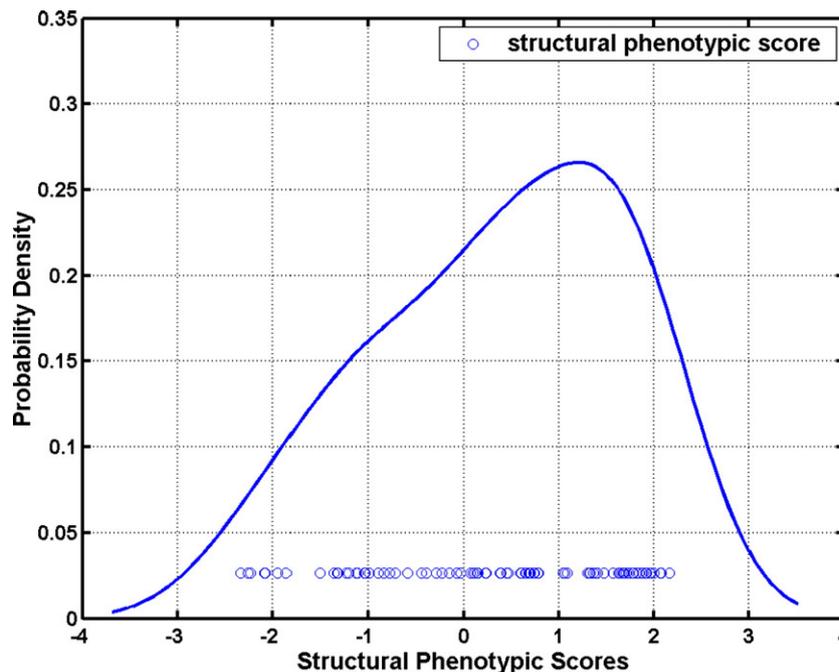


Fig. 6. Histograms of the MRI-based classification scores for MCI subjects obtained via applying the classifiers built on AD and CN participants. Fifty-seven out of 88 MCI subjects display positive scores, i.e. their MRI scans indicate that they possess the structural pattern characteristic of AD.

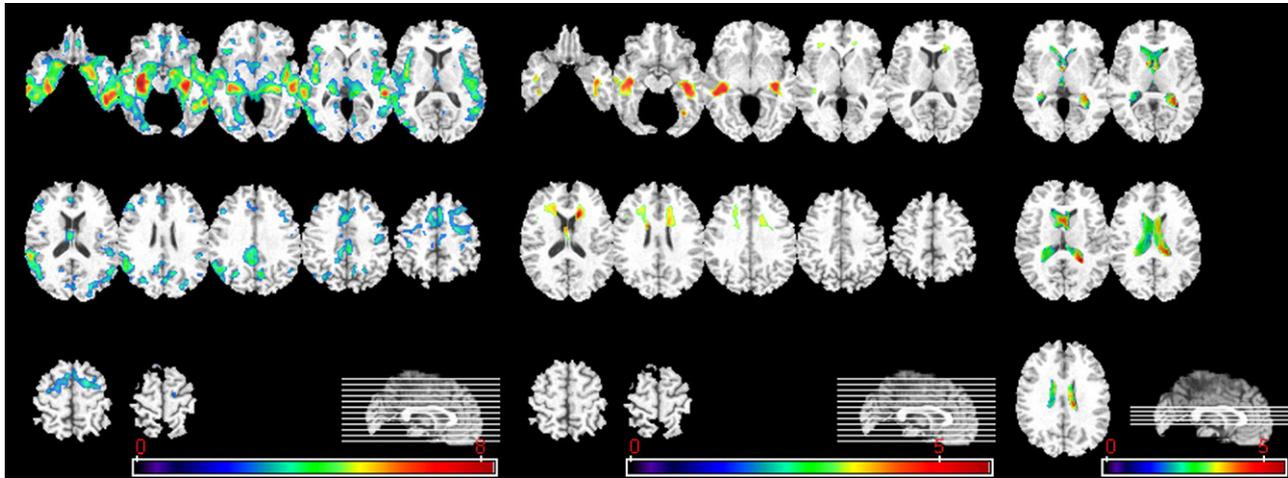


Fig. 7. Voxel-based analysis of group differences between MCL_{CN} and MCL_{AD} . From left to right, group comparison results on GM, WM, and CSF are shown (GM, WM: $MCL_{CN} > MCL_{AD}$, CSF: $MCL_{AD} > MCL_{CN}$, $p < 0.05$, corrected). The color-maps indicate the scale for the t -statistic. Images are displayed in radiological convention.

validation, which were obtained from these two subgroups are shown in Fig. 12, for different values of the threshold on MMSE rate of change. These results indicate that maximal separation between MCI_{PR} and MCI_{NPR} is obtained for a threshold close to -1 , which is in agreement with our expectations. The classification rate obtained at that threshold was 87% and the area under the curve was 0.86.

Discussions and conclusion

This study utilized computational neuroanatomic methods to quantify cross-sectional patterns of brain atrophy in a relatively large sample of cognitively normal elderly individuals, and in patients with MCI and AD. Spatially complex spatial patterns of brain atrophy were measured, and were found to be consistent with known patterns of AD pathology from histological studies. MCI patients had significant temporal lobe atrophy, especially in the hippocampus, superior, inferior temporal gyrus, and uncus, as well

as medial GM atrophy, especially in the posterior cingulate and adjacent precuneus, and the medial aspect of the uncus. Additional GM atrophy was measured between MCI and AD patients, particularly in the hippocampus, the entorhinal cortex, and the middle and inferior temporal gyrus. The pattern of atrophy also included the WM surrounding the hippocampus, and the ventricles, albeit at much lower significance, compared to GM atrophy.

The complexity of this pattern of atrophy suggests that perhaps more sophisticated methods for measuring structural brain changes in MCI and AD can be helpful for diagnosis and prognosis of the disease, compared to the most common approach that has been taken up to date in the neuroimaging literature (Kaye et al., 1997; Convit et al., 2000; Killiany et al., 2000; Dickerson et al., 2001; Chetelat et al., 2002), namely to examine volumes of a small number of structures typically of the hippocampus and the entorhinal cortex. This is further bolstered by histopathological studies (Braak et al., 1998) that have investigated the pattern of deposition of β -amyloid plaques and tau-pathology during the

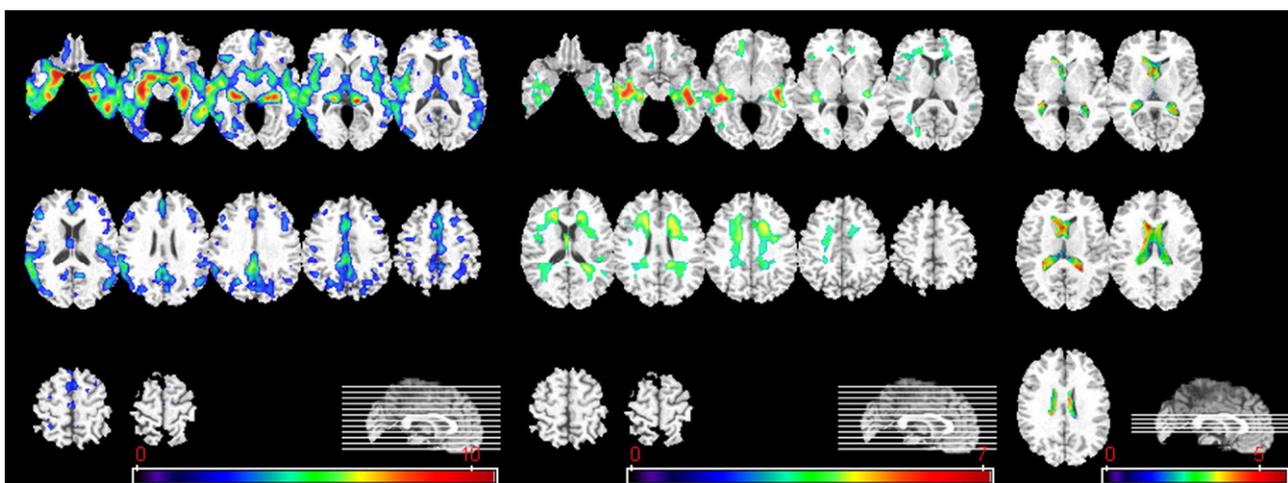


Fig. 8. Voxel-based analysis of group difference between CN and MCL_{AD} . From left to right, group comparison results on GM, WM, and CSF are shown (GM, WM: $CN > MCL_{AD}$, CSF: $MCL_{AD} > CN$, $p < 0.05$, corrected). The color-maps indicate the scale for the t -statistic. Images are displayed in radiological convention.

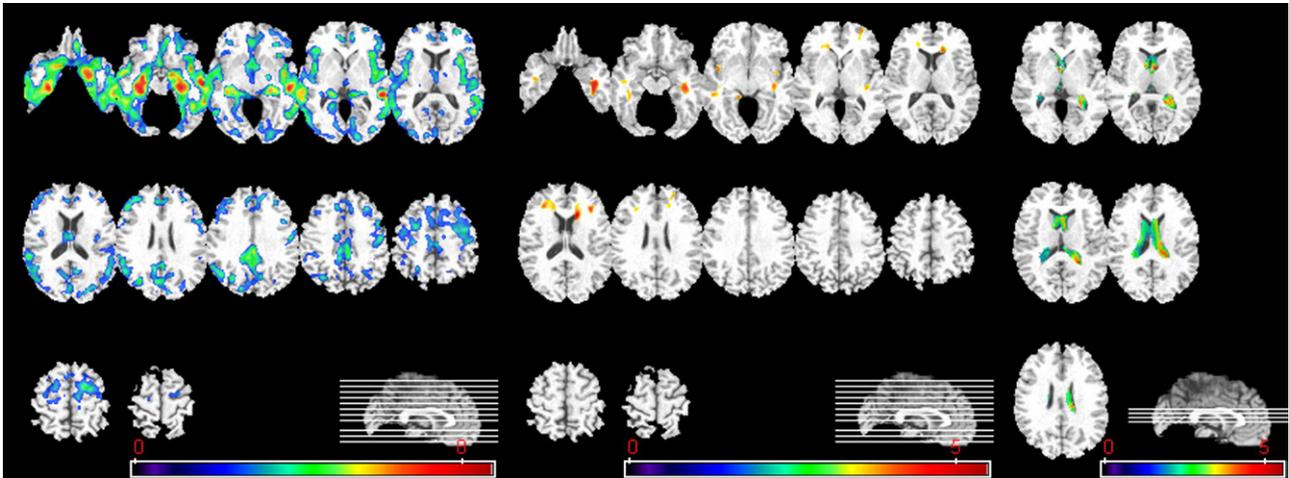


Fig. 9. Voxel-based analysis of group difference between MCL_{CN} and AD. From left to right, group comparison results on GM, WM, and CSF are shown (GM, WM: MCL_{CN}>AD, CSF: AD>MCL_{CN}, $p<0.05$, corrected). The color-maps indicate the scale for the t -statistic. Images are displayed in radiological convention.

progression of AD, as well as with studies of magnetization transfer that indicated a more than expected widespread distribution of brain pathology (Van der Flier et al., 2002). The results of the current study also demonstrated that sole measurements of the hippocampus and the entorhinal cortex are not sufficient for separating the three groups from each other, not even AD from CN, with clinically adequate sensitivity and specificity, since hippocampal and entorhinal cortex measurements were highly overlapping between MCI and AD or CN (Fig. 1). Overlap was also observed between AD and CN.

Perhaps the most exciting finding of the current study is that the MCI subgroup identified by the classifier as AD-like showed a markedly faster rate of subsequent MMSE decline, whereas the group that had similar SPS to CN showed minor MMSE decline.

Related was the converse finding, namely that MCI individuals whose MMSE scores decreased relatively more rapidly were relatively well distinguishable from the ones that show no decline or relatively small decline similar to that of CN individuals. This finding indicates that the SPS score determined through pattern analysis and classification has predictive clinical value, which would render it a significant biomarker for early AD stages and for individuals that are good candidates for treatment. Importantly, the SPS scores were derived using a single cross-sectional MRI scan, and not from longitudinal scans, which renders them more practically feasible from a logistical as well as from a financial point of view.

One of the main conclusions of this study is that two thirds of the MCI patients of this cohort are closer to AD than they are to

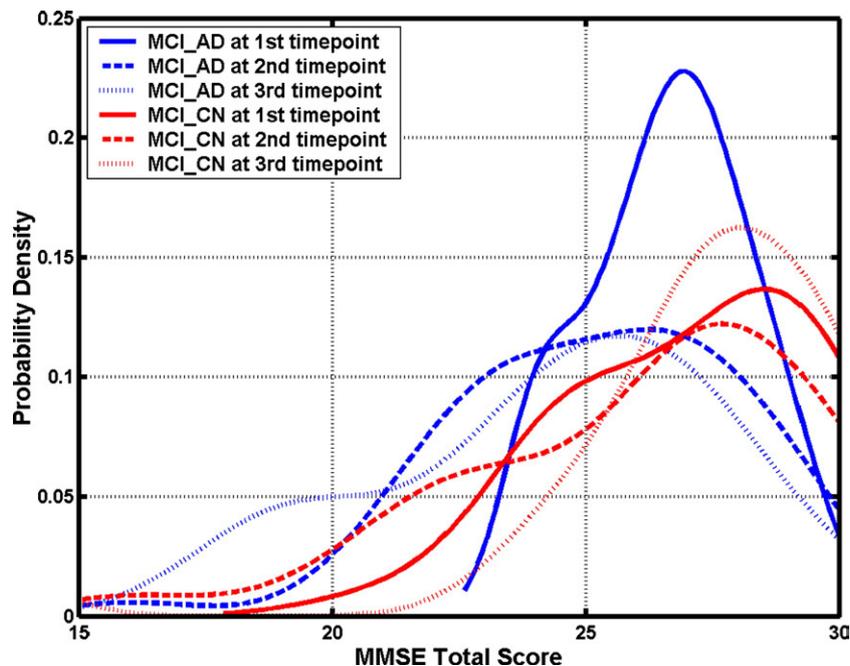


Fig. 10. The distributions of the MMSE scores during baseline and follow-ups of MCI subgroups.

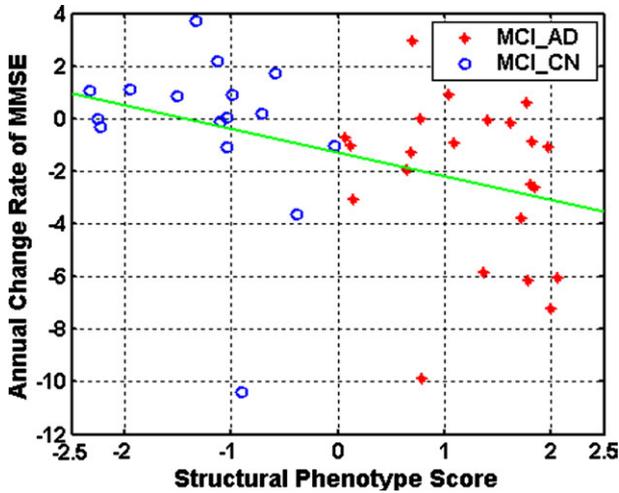


Fig. 11. Regression plot of the rates of MMSE change and the SPS scores at baseline.

cognitively normal individuals. Although previous MRI-based studies of MCI patients have demonstrated brain atrophy mainly in medial temporal lobe structures (Convit et al., 1997; Jack et al., 1999; Jack et al., 2000; Xu et al., 2000; De Santi et al., 2001; Du et al., 2001; Grundman et al., 2002; Chetelat, 2003; Karas et al., 2004; Pennanen et al., 2005; Bozzali et al., 2006; Saykin et al., 2006; Whitwell et al., 2007), the present study finds that brain atrophy in this cohort is already quite extensive and involves superior, middle, and inferior temporal gyri, the insula, the

posterior cingulate and adjacent precuneus, the uncus, and the peri-hippocampal WM, mainly in the MCI_{AD} subgroup. Two factors might have contributed to this finding. First, the ADNI MCI population was selected so that an adequate number of converters would be expected within 3 years, so that the study would be adequately powered. Although our findings certainly agree with this clinically based selection of patients, Fig. 1 suggests that the volumes of the hippocampus and entorhinal cortex of the MCI patients were highly overlapping with both CN and AD; the cross-validated classification results obtained from these ROI measurements also showed significant group overlap. Therefore, patient selection does not fully explain our finding. It is likely that the identification of more widespread and complex patterns of brain atrophy in our study is partly due to the fact that a high-dimensional template warping mechanism was used to determine the RAVENS maps and to capture spatial patterns of brain atrophy. This image warping algorithm has been previously found to achieve very accurate inter-individual registration, which is of fundamental importance for measuring subtle patterns of brain atrophy across individuals.

The similarity of a subgroup of the MCI group to AD was further supported by the complementary analysis using high-dimensional pattern classification to determine the optimal group separation. This analysis showed that the structural phenotypic scores of two thirds of the MCI group were more similar to those of AD patients. Relatively recent studies using the PIB compound have also shown a relatively widespread accumulation of amyloid plaques in many MCI patients (Kempainen et al., 2007). These findings further support that AD pathology might already be at quite advanced stages by the time cognitive decline becomes

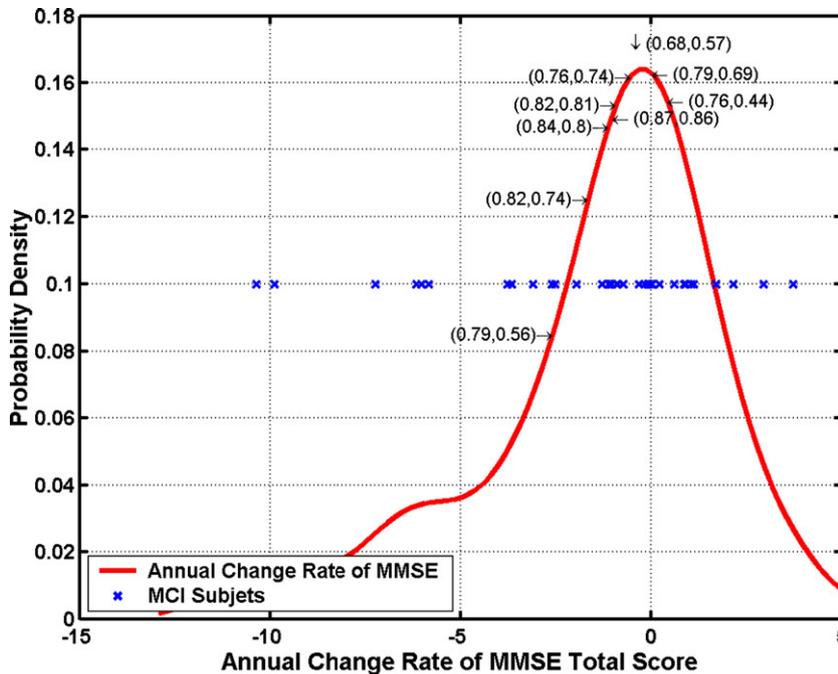


Fig. 12. Classification rates and areas under the ROC curve obtained by subgrouping MCI patients into progressors and nonprogressors, according to a threshold on their rates of MMSE change within a year. Optimal classification rate of 0.87=87% (AUC 0.86) was obtained for a threshold around -1, i.e. if one defines progressors as the MCI patients that display rates of change of MMSE score < -1/year. This is in agreement with the fact that even CN individuals display some rate of decline. The red curve is a histogram of the rate of MMSE change, the blue stars are individual MCI patients, and the numbers before and after commas within parentheses are the correct classification rates and the AUCs.

clinically detectable, at least for a subgroup of the MCI population that is hypothesized to have significant AD pathology. Subsequent follow-up will determine whether this hypothesis is true; however, the MMSE decline in this subgroup is a significant indicator that these patients are likely to convert to AD soon. The ability of pattern classification to serve as a biomarker for such a group would be very important.

The finding that the majority of MCI patients seem to have AD-like structural profiles also suggests that more emphasis should be placed on studying CN groups. Although current clinical trials of potential treatments, including ADNI, focus primarily on MCI groups, since MCI patients convert to AD at rates of approximately 15% annually, from a diagnostic perspective it would undoubtedly be beneficial to study cognitively normal populations that have less advanced AD pathology. By virtue of its ability to measure subtle patterns of brain atrophy, the methodology adopted in our study can potentially assist in identifying cognitively normal individuals that display patterns of atrophy that render them likely to be in a very early preclinical AD stage. Earlier analysis using the same methodology in a longitudinal study of normal aging demonstrated that high-dimensional pattern analysis and classification can identify abnormal patterns of brain atrophy before clinically detectable cognitive decline (Davatzikos et al., 2006).

The finding of reduced WM volumes between MCI and CN is interesting and merits further research. The pattern of WM atrophy was bilateral, although more pronounced in the right hemisphere, and extended into the region adjacent to the entorhinal cortex as well as into the superior and middle temporal gyri. Dense connections existing between the hippocampus and the posterior cingulate, which coupled with the early changes that have been reported in the posterior cingulate (Chetelat, 2003; Chetelat et al., 2003), might imply that changes in WM might provide additional markers of disease progression, something that has traditionally not attracted much attention in the AD literature. A growing recent literature using diffusion tensor imaging further supports the importance of examining white matter changes in AD (Bozzali et al., 2002; Moseley, 2002; Fellgiebel et al., 2004; Choi et al., 2005; Fellgiebel et al., 2005, 2006; Medina et al., 2006; Naggara et al., 2006; Ray et al., 2006; Huang and Auchs, 2007), albeit the majority of these studies have been restricted to measuring quantities such as fractional anisotropy and diffusivity, and therefore have not differentiated between brain atrophy and other tissue changes that can potentially have vascular underpinnings (for example, both fractional anisotropy and diffusivity are known to be lower in leukoencephalopathy). More sophisticated types of analysis of diffusion tensor images (Khurd et al., 2006; Verma and Davatzikos, 2006) can potentially elucidate alterations of WM connectivity in AD.

Our findings suggest a bilateral pattern of atrophy in MCI, although the right hemisphere displayed higher magnitude and more widespread extent of atrophy of both GM and WM. This potential asymmetry was, however, balanced with disease progression, since the pattern of atrophy in AD was fairly symmetric. The interpretation of such asymmetries is known to be problematic, since the true reason might be bias in patient selection rather than differences of the underlying AD pathology. In particular, the right-more-than-left pattern that we observed is consistent with the hypothesis that patients that report to the clinic with memory complaints are more likely to report when they have language problems. Accordingly, one might expect that a smaller degree of atrophy on the left hemisphere would meet the threshold

for a patient's reporting to the clinic, compared to atrophy of the right hemisphere that would be likely to present less obvious cognitive deficits. Put differently, a relatively larger degree of right-hemisphere atrophy, compared to left, is likely to be tolerated before the patient reports to the clinic. Our pattern of asymmetry is the reverse of what another similar study reported (Karas et al., 2004). Differences between the two studies, especially with respect to the template warping method and the patient populations, render the two studies not directly comparable. The relatively higher sensitivity of our methodology in detecting GM and WM atrophy (e.g. values of the *t*-statistic in Figs. 2 and 3) further speaks to the methodological differences between the two studies.

The comparison between the MCI_{CN} and MCI_{AD} subgroups leads to two very interesting conclusions. First, the former group is almost entirely overlapping with CN, and the latter overlaps almost entirely with the AD group. Although this result yet remains to be tested in independent patient populations, it does highlight the potential of the high-dimensional pattern classification method to detect subgroups in MCI patients, which would be of great importance clinically. Second, the main WM differences between these two subgroups were in periventricular tissue. This finding could imply decrease of WM via Wallerian degeneration; however, testing this hypothesis would require different imaging protocols, and especially diffusion tensor imaging, which is not available in ADNI. It is interesting to note, however, that the regions identified by this analysis are exactly where the bulk of leukoencephalopathy tends to occur in elderly individuals with or without other significant vascular disease, and which tend to appear as gray rather than white matter in T1-weighted MR images. The resolution and contrast of the MRI sequences used in this study do not allow us to investigate this issue. This finding raises the important issue of the potential role of vascular pathology in AD, which has also received attention in the literature (Snowdon et al., 1997; Schneider et al., 2003; Prins et al., 2004; Schneider et al., 2004; Kim et al., 1998; Lin et al., 1999; Bennett et al., 2000; Shi et al., 2000; Nihashi et al., 2001). Our results suggest that one of the significant differences between the MCI_{CN} and MCI_{AD} subgroups is likely to be periventricular leukoencephalopathy, and further support the need to examine vascular pathology in tandem with brain atrophy. Regardless of whether or not AD is pathophysiologically related to vascular disease, its clinical manifestation almost certainly depends on the concurrent presence of vascular disease (Schneider et al., 2004).

A relatively new technique, namely high-dimensional pattern classification, was used to analyze patterns of spatial distribution of brain tissue and integrate them into an abnormality score, which represents how similar the entire structural profile of an individual fits that of AD patients or of cognitively normal individuals. This approach has been recently used in several neuroimaging studies and has shown great potential as a diagnostic tool on individuals (Lao et al., 2004; Davatzikos et al., 2005a,b; Fan et al., 2007; Davatzikos et al., in press). It is a significant deviation from either ROI-based or voxel-based techniques, which examine the brain region-by-region independently, without integrating the entire pattern of atrophy (or functional activity; Davatzikos et al., 2005a,b) throughout all brain regions together. This is very important, because although many regions generally display significant group differences, they also significantly overlap between groups (see Fig. 1), and therefore do not offer sufficient sensitivity and specificity for diagnostic purposes. The methodology used herein achieves high group separation via nonlinear multivariate

classification using support vector machines, and it has been shown to possess great diagnostic value in neurodegenerative and neuropsychiatric disorders, and likely beyond (Zhang et al., 2002; Davatzikos et al., 2005a,b; LaConte et al., 2005; Mourao-Miranda et al., 2005).

In summary, this study used advanced quantitative pattern analysis and classification methodologies and determined spatially complex patterns of brain atrophy in MCI and AD patients. The MCI group was highly variable, as anticipated, but its majority overlapped with AD patients, with regard to brain atrophy. Analysis of the follow-up scans of this longitudinal study revealed that the group identified by pattern classification as being similar to the AD group did indeed present significantly higher rates of MMSE decline. Further follow-up will help reveal whether subsequent conversion of the MCI participants to AD will be in agreement with the two structural profiles observed in this study, i.e. whether the MCI subgroup that overlaps with AD will indeed progress to AD within the 3-year follow-up period of ADNI.

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