

Region-Specific Hierarchy between Atrophy, Hypometabolism, and β -Amyloid ($A\beta$) Load in Alzheimer's Disease Dementia

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Gray matter atrophy, glucose hypometabolism, and β -amyloid $A\beta$ deposition are well-described hallmarks of Alzheimer's disease, but their relationships are poorly understood. The present study aims to compare the local levels of these three alterations in humans with Alzheimer's disease. Structural magnetic resonance imaging, ¹⁸F-fluorodeoxyglucose positron emission tomography (PET), and ¹⁸F-florbetapir PET data from 34 amyloid-negative healthy controls and 20 demented patients with a high probability of Alzheimer's disease etiology (attested using neuroimaging biomarkers as recently recommended) were analyzed. For each patient and imaging modality, age-adjusted Z-score maps were computed, and direct between-modality voxelwise comparison and correlation analyses were performed. Significant differences in the levels of atrophy, hypometabolism, and $A\beta$ deposition were found in most brain areas, but the hierarchy differed across regions. A cluster analysis revealed distinct subsets of regions: (1) in the hippocampus, atrophy exceeded hypometabolism, whereas $A\beta$ load was minimal; (2) in posterior association areas, $A\beta$ deposition was predominant, together with high hypometabolism and lower but still significant atrophy; and (3) in frontal regions, $A\beta$ deposition was maximal, whereas structural and metabolic alterations were low. Atrophy and hypometabolism significantly correlated in the hippocampus and temporo-parietal cortex, whereas $A\beta$ load was not significantly related to either atrophy or hypometabolism. These findings provide direct evidence for regional variations in the hierarchy and relationships between $A\beta$ load, hypometabolism, and atrophy. Altogether, these variations probably reflect the differential involvement of region-specific pathological or protective mechanisms, such as the presence of neurofibrillary tangles, disconnection, as well as compensation processes.

Introduction

Alzheimer's disease is characterized by major brain changes, including gray matter atrophy, hypometabolism, and β -amyloid ($A\beta$) deposition. These changes have been described extensively using neuroimaging techniques, i.e., magnetic resonance imaging (MRI)

and positron emission tomography (PET) combined with specific radiotracers: ¹⁸F-fluorodeoxyglucose (FDG) for glucose metabolism, ¹¹C-Pittsburgh compound B and ¹⁸F-florbetapir or others for $A\beta$ deposition (Herholz and Ebmeier, 2011). These techniques have proved to be useful for early diagnosis of Alzheimer's disease but also for additional understanding the pathological mechanism(s) underlying the disease (Rabinovici and Roberson, 2010).

Dynamic models of changes in these neuroimaging biomarkers over the course of the disease have been proposed recently (Perrin et al., 2009; Jack et al., 2010; Ewers et al., 2011). According to the amyloid cascade hypothesis (Hardy and Selkoe, 2002), the sequence of events would start with $A\beta$ deposition in a very early, presymptomatic stage. Then, $A\beta$ is thought to trigger functional and structural changes that would appear later, closer to clinical symptom onset. However, despite experimental studies showing that $A\beta$ impairs synaptic functioning (Palop and Mucke, 2010; Parihar and Brewer, 2010), results from the human neuroimaging literature are ambiguous. Correlations between increased $A\beta$

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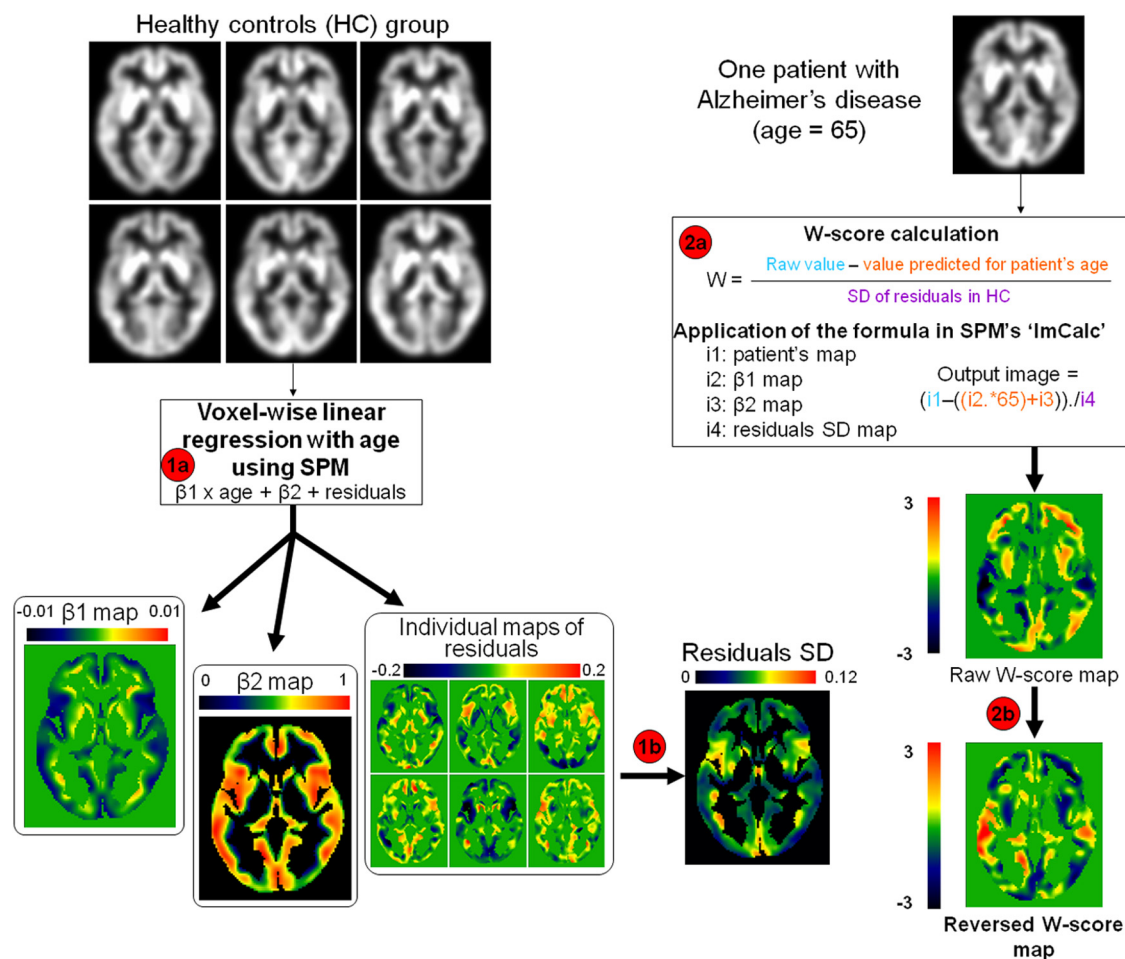


Figure 1. Illustration of the different steps required to create *W*-score maps using the SPM software. This figure shows the procedure for the atrophy *W*-score map of a 65-year-old patient with Alzheimer's disease compared with six healthy controls (HC) for the sake of illustration. First, a simple linear regression was performed in the HC group to estimate age-related changes (**1a**), resulting in several files: the β_1 map containing voxelwise age-related regression coefficients, the β_2 map containing intercept values, and the individual maps of residuals. The SD of residuals was computed voxelwise (**1b**). A *W*-score map was then created using the corresponding formula and previously computed maps using the SPM "ImCalc" function (**2a**). Last, for MRI and FDG–PET data, *W* values were reversed so that positive numbers represent pathological features in all three imaging modalities (**2b**).

and decreased metabolism in Alzheimer's disease patients have been found in some studies (Engler et al., 2006; Edison et al., 2007; Cohen et al., 2009) but not in others (Li et al., 2008; Rabinovici et al., 2010; Furst et al., 2012). Previous works showed that the relationship between $A\beta$ load and hypometabolism (Cohen et al., 2009) or atrophy (Chételat et al., 2010) varies throughout the evolution of the disease, suggesting that neurodegeneration could be independent of $A\beta$ pathology in later stages (Hyman, 2011).

Moreover, there may also be differences in the sequence of events according to brain regions, as suggested by discrepancies in the regional pattern of atrophy, hypometabolism, and $A\beta$ deposition. More specifically, although all three alterations are found in some regions, such as the posterior cingulate, precuneus, and temporo-parietal areas (Buckner et al., 2005; Jack et al., 2008), regional discrepancies have been highlighted when comparing modalities two by two. Thus, differential degrees of atrophy and hypometabolism are found in the posterior cingulate cortex versus the hippocampus, for example (Alsop et al., 2008; Chételat et al., 2008), and both atrophy and hypometabolism can be found in regions with low $A\beta$ load or can be absent in regions with high $A\beta$ deposition (Edison et al., 2007; Jack et al., 2008; Li et al., 2008).

The main objective of this study was therefore to characterize and compare the regional degrees of gray matter atrophy, hypometabolism, and $A\beta$ deposition as measured in the same patients with Alzheimer's disease. In addition, we aimed at assessing for the first time the local correlations between these three alterations using a voxel-based method. For these purposes, we thoroughly selected patients as having a high probability of Alzheimer's disease etiology according to recent recommendations (McKhann et al., 2011) and used a methodology especially designed to compare data from different imaging modalities (Chételat et al., 2008).

Materials and Methods

Participants

Twenty-two patients diagnosed with probable Alzheimer's disease were first selected according to National Institute of Neurological and Communicative Disorders and Stroke/Alzheimer's Disease and Related Disorders Association clinical criteria (McKhann et al., 1984). Following the recent recommendations from the National Institute on Aging and Alzheimer's Association workgroup (McKhann et al., 2011) for research studies, only those 20 patients with a high probability of Alzheimer's disease etiology (i.e., who have positive neuroimaging biomarkers for both $A\beta$ deposition and neurodegeneration) were selected for additional analyses (see below).

Table 1. Selection of patients with a high probability of Alzheimer's disease etiology based on neuroimaging biomarkers of both amyloid deposition and neuronal degeneration (McKhann et al., 2011)

Type of biomarker	Index used	Patients with a clinical diagnosis of probable Alzheimer's disease																					
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22
A β deposition	Florbetapir neocortical SUVR	1.46	1.54	1.86	1.20	1.48	1.81	0.87	1.80	1.92	1.71	1.41	1.77	1.56	1.28	1.29	1.83	1.72	0.89	1.59	1.68	1.45	1.76
Neuronal degeneration	Atrophy <i>W</i> -score	3.36	1.82	5.71	1.33	3.52	4.77	6.11	3.24	2.37	2.07	1.36	2.51	0.13	1.08	2.84	1.91	4.25	2.01	1.72	2.85	3.80	1.38
	Hypometabolism <i>W</i> -score	1.39	4.44	4.88	1.75	5.51	5.83	1.96	3.26	3.26	4.33	2.97	4.42	1.73	1.89	3.05	5.52	4.83	2.30	2.40	4.84	1.62	4.30
Presence of both biomarkers?		Yes	Yes	Yes	Yes	Yes	Yes	No	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	No	Yes	Yes	Yes	Yes

Amyloid load was assessed through neocortical florbetapir SUVR and considered as positive over 1.1 (for more details, see Materials and Methods). Alzheimer's disease type neuronal degeneration was considered using two indexes: (1) atrophy in the hippocampus, amygdala, and lateral temporal lobe and (2) hypometabolism in the angular gyrus, precuneus, and posterior cingulate. Both indexes are expressed in *W*-scores (i.e., age-adjusted *Z*-scores) compared with the control group. Patients were selected for additional analysis if they had a florbetapir-positive PET scan together with at least one neurodegeneration *W*-score > 1.65. Indexes considered as positive are bolded. This condition was fulfilled for all patients, except for patients 7 and 18 that were consequently excluded from additional analyses.

Thirty-seven healthy controls were also enrolled in this study after clinical and neuropsychological examination. They had no history or clinical evidence of major neurological or psychiatric disorder and performed in the normal range in all neuropsychological tests (including tests of episodic memory, working memory, language skills, executive functions, and visuospatial abilities).

The study was approved by the local ethics committee, and all participants gave written consent for participation before the scans.

Neuroimaging data acquisition

All participants were scanned on the same MRI and PET cameras at the Cycleron Centre (Caen, France). The median time lapse between the first and last examination was 16.5 d (minimum of 1; maximum of 60) for patients and 28 d (minimum of 8; maximum of 215) for controls.

MRI data. For each participant, a high-resolution T1-weighted anatomical image was acquired on a Philips Achieva 3 T scanner using a three-dimensional fast-field echo sequence (sagittal; repetition time, 20 ms; echo time, 4.6 ms; flip angle, 20°; 170 slices; slice thickness, 1 mm; field of view, 256 × 256 mm²; matrix, 256 × 256).

PET data. Both FDG and florbetapir PET scans were acquired on a Discovery RX VCT 64 PET–CT device (GE Healthcare) with a resolution of 3.76 × 3.76 × 4.9 mm (field of view, 157 mm). Forty-seven planes were obtained with a voxel size of 2.7 × 2.7 × 3.27 mm. A transmission scan was performed for attenuation correction before the PET acquisition.

FDG–PET. Participants were fasted for at least 6 h before scanning. After a 30 min resting period in a quiet and dark environment, ~180 MBq of FDG was intravenously injected as a bolus. A 10 min PET acquisition scan began 50 min after injection.

Florbetapir–PET. Each participant underwent a 20 min PET scan, beginning 50 min after the intravenous injection of ~4 MBq/kg of florbetapir.

Neuroimaging data handling and transformation

Preprocessing. MRI data were segmented, normalized, and modulated using the VBM5.1 toolbox (<http://dbm.neuro.uni-jena.de>), implemented in the Statistical Parametric Mapping 5 (SPM) software (Wellcome Trust Centre for Neuroimaging, London, UK) to obtain maps of local gray matter volume corrected for brain size. PET data (both FDG and florbetapir) were corrected for partial volume effects (PMOD Technologies), coregistered onto their corresponding MRI, and normalized using the deformation parameters defined from the MRI procedure. Resultant images underwent quantitative scaling using the cerebellar gray matter as a reference to obtain standardized uptake value ratio (SUVR) images.

Because MRI and PET data have different original spatial resolutions, a differential smoothing was applied to equalize the effective smoothness (Richardson et al., 1997; Chételat et al., 2008; Villain et al., 2008): a Gaussian kernel of 10 × 10 × 10 (*x*, *y*, *z*) mm was used for the MRI data and 9.3 × 9.3 × 8.8 mm for the PET data. Finally, images were masked to exclude non-gray matter voxels from the analyses.

Creation of *W*-score maps. To obtain measurements of atrophy, hypometabolism, and amyloid load expressed in the same unit, therefore

Table 2. Demography and mini-mental state examination scores in the groups used for data analyses

	AD (<i>n</i> = 20)	HC (<i>n</i> = 34)	<i>p</i> value
Age (years)	68.9 ± 9.0	68.1 ± 7.2	0.83
Women (%)	10 (50.0%)	20 (58.8%)	0.73
Education (years)	10.35 ± 3.8	11.8 ± 3.7	0.11
MMSE	20.6 ± 4.5	29.1 ± 0.8	<0.001

Statistical analyses were performed using Mann–Whitney *U* test [for age, education, and mini-mental state examination (MMSE)] and Yates's χ^2 test (sex ratio). AD, Alzheimer's disease patients; HC, healthy controls.

enabling a direct comparison of different imaging modalities, *W*-score maps were computed for each patient and each imaging modality using the healthy control group as a reference (Fig. 1). *W*-scores are analogous to *Z*-scores but they are adjusted for specific covariate(s) (Jack et al., 1997; Boccardi et al., 2003; Jack et al., 2008), age in the present case. Like *Z*-scores, *W*-scores have a mean value of 0 and an SD of 1 in the control group, and values of +1.65 and −1.65 correspond to the 95th and 5th percentiles, respectively. To create *W*-score maps, voxelwise regressions were first performed in the control group between age and each imaging data using the SPM software. Then, *W*-score maps were computed using the following formula: *W*-score = [(patient's raw value) − (value expected in the control group for the patient's age)]/SD of the residuals in controls.

For MRI and FDG–PET, *W*-scores were reversed so that positive *W*-scores indicate pathology for all modalities (i.e., less gray matter volume, less glucose metabolism, more A β deposition). For each modality, individual *W*-score maps were averaged across the patients to provide whole-brain profiles of atrophy, hypometabolism, and A β deposition, expressed as mean *W*-scores.

Regions of interest. Finally, to obtain quantitative and statistical information in specific brain areas, mean values of gray matter volume, FDG SUVR, and florbetapir SUVR were extracted from the corresponding images before the smoothing step in 11 regions of interest defined using the automated anatomical labeling atlas (Tzourio-Mazoyer et al., 2002): hippocampus, amygdala, parahippocampus, temporal pole, angular gyrus, precuneus/posterior cingulate, lateral temporal, anterior cingulate, orbitofrontal, dorsomedial prefrontal, and dorsolateral prefrontal cortices. The neocortical values of florbetapir retention were also extracted from the florbetapir SUVR images for the sake of classification of patients as amyloid positive versus negative (see below).

Selection of patients and subjects using neuroimaging biomarkers

Patients. As mentioned above, neuroimaging biomarkers were used to further select those Alzheimer's disease patients with a high likelihood of Alzheimer's disease etiology, i.e., showing positive biomarkers of both A β deposition (from florbetapir–PET) and neurodegeneration (from MRI or FDG–PET), as recommended (McKhann et al., 2011). First, values of florbetapir neocortical SUVR were used to separate amyloid-positive from amyloid-negative patients. For that purpose, a florbetapir SUVR cutoff value was determined using a cluster analysis performed on

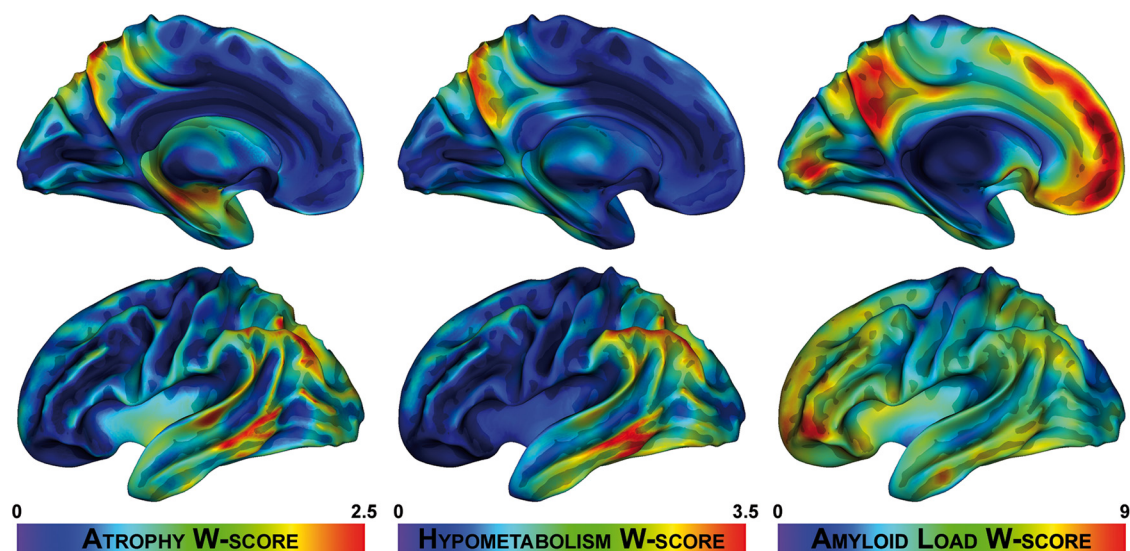


Figure 2. Brain patterns of alteration in the 20 patients with Alzheimer's disease dementia. For each imaging modality, local degrees of alteration are expressed as mean W -score compared with the control group ($n = 34$) in each gray matter voxel. Note that, for all imaging modalities, a positive W -score indicates a pathological feature. Colors have been scaled to the range of each modality to fit to the regional distribution of each process. For clarity, only the left hemisphere is represented here, because results were mainly symmetrical.

the whole sample (Bourgeat et al., 2010; Rowe et al., 2010), yielding to a value of 1.1. This value is consistent with previous studies using florbetapir (Fleisher et al., 2011; Camus et al., 2012).

Then, neurodegeneration was assessed using two specific indexes. For atrophy, the hippocampus, amygdala, and lateral temporal cortex were combined, and a global W -score was computed. Hypometabolism W -score included the angular gyrus, precuneus, and posterior cingulate. These areas were chosen as the prototypical neural substrates of Alzheimer's disease, according to a recent meta-analysis of MRI and FDG–PET studies (Schroeter et al., 2009).

Patients were included if they had a positive florbetapir–PET scan and at least one positive biomarker of neurodegeneration (W -score ≥ 1.65 for atrophy and/or hypometabolism). Individual values for the three biomarkers are shown in Table 1. Of the 22 clinically diagnosed patients, two showed a negative florbetapir–PET scan. For the 20 remaining patients, at least one of the two biomarkers of neuronal injury was positive. Consequently, these 20 Alzheimer's disease patients were all included in the following analyses. Only three of these 20 patients were treated with cholinesterase inhibitors when included, whereas the others were *de novo* (untreated) patients. All analyses were repeated without these three patients, and the conclusions remained unchanged (data not shown) so that all results will be presented on the complete sample of 20 patients.

Healthy controls. In addition, to avoid the presence of healthy subjects at a preclinical stage of Alzheimer's disease in the control group that could bias our analyses, amyloid-positive subjects were excluded from additional analysis. Indeed, the presence of cerebral amyloidosis is believed to indicate the first stage of preclinical Alzheimer's disease (Sperling et al., 2011). Therefore, three subjects with a florbetapir SUVR value > 1.1 were withdrawn from the control group.

Demographic data and mini-mental state examination scores for the 34 controls (14 males, 20 females) and 20 patients with highly probable Alzheimer's disease (10 males, 10 females) are displayed in Table 2.

Statistical analyses

Comparing local atrophy, hypometabolism, and $A\beta$ burden: voxelwise analyses. Individual W -score maps of atrophy, hypometabolism, and $A\beta$ deposition were compared in a voxelwise factorial analysis using the SPM software, with MRI, FDG–PET, and florbetapir–PET images as within-subject measurements. Results were considered as significant at p [familywise error (FWE) corrected for multiple comparisons] < 0.05 threshold and cluster size $k \geq 20$ (160 mm³).

Regions of interest analyses. Using the values extracted from the 11 regions of interest, regional W -scores were calculated and compared us-

ing nonparametric statistical tests. To compare the local degrees of atrophy, hypometabolism, and $A\beta$ deposition, a Friedman's ANOVA was performed within each region of interest. If significant (i.e., $p < 0.05$), *post hoc* two-by-two comparisons were conducted using Wilcoxon's signed-rank test. So as to provide a more global insight on the different gradients of alteration across brain regions, a Ward's hierarchical clustering method was used to group brain regions with closely similar patterns.

Correlation between local alterations. To assess the link between local atrophy, hypometabolism, and $A\beta$ deposition, W -score maps were entered two by two in voxelwise correlation analyses using the Biological Parametric Mapping toolbox implemented in SPM and especially designed to analyze brain images from different modalities (Casanova et al., 2007). Three independent correlation analyses were performed: $W_{A\beta}$ versus $W_{\text{hypometabolism}}$, $W_{A\beta}$ versus W_{atrophy} , and W_{atrophy} versus $W_{\text{hypometabolism}}$. Results were considered as significant at a $p_{\text{FWE}} < 0.05$ threshold and cluster size $k \geq 20$ (160 mm³).

Results

Patterns of gray matter atrophy, hypometabolism, and $A\beta$ deposition in Alzheimer's disease

Averaged W -score maps for each modality are shown in Figure 2. Briefly, gray matter atrophy was found in the medial and lateral temporal, inferior parietal cortex, and precuneus. Hypometabolism concerned the precuneus, posterior cingulate, lateral temporal and parietal cortices, and, to a lesser degree, the medial temporal lobe. $A\beta$ deposition was found in most brain areas, with highest W -scores in the medial and orbital prefrontal cortex, precuneus, and posterior cingulate (with W -score ≥ 9), whereas the primary sensorimotor cortex, occipital cortex, thalamus, and medial temporal lobe were relatively spared.

Voxelwise comparisons between alterations

The results of the direct between-modality comparison analyses are displayed in Figure 3. First, the comparison between atrophy and hypometabolism revealed significant differences in both directions. Hypometabolism was greater than atrophy in the precuneus, posterior cingulate, parietal, and middle/inferior temporal gyrus, whereas atrophy significantly exceeded hypometabolism in the anterior part of medial temporal lobe. Second, $A\beta$

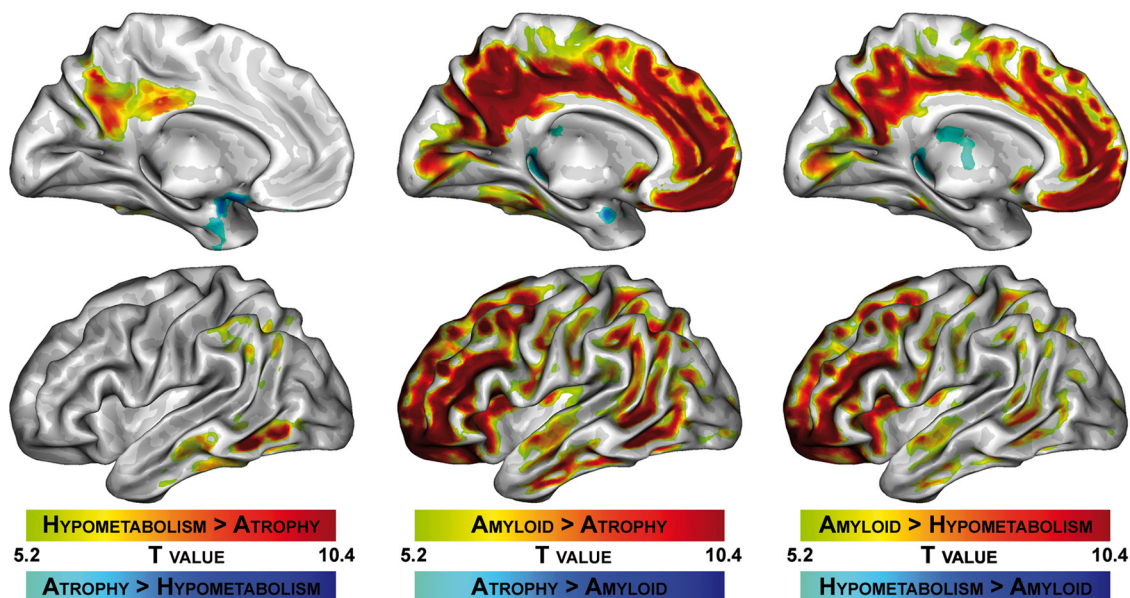


Figure 3. Voxelwise comparisons between the local degrees of atrophy, hypometabolism, and A β deposition in the 20 patients with Alzheimer's disease dementia. The t value of 5.2 used as a threshold in this figure corresponds to the p (FWE corrected) < 0.05 threshold described in Results.

deposition significantly exceeded atrophy in most brain areas, with greatest differences in frontal areas, precuneus, and posterior cingulate and less but still significant differences in lateral temporal and parietal regions. Conversely, atrophy was superior to A β deposition in the anterior part of the medial temporal lobe and in the posterior hippocampus. Third, A β deposition considerably exceeded hypometabolism in prefrontal cortex, precuneus, and posterior cingulate, whereas the difference was less marked, but still significant, in the lateral temporal and parietal cortices. Finally, hypometabolism significantly exceeded A β deposition in the thalamus and posterior hippocampus.

Voxelwise correlations between alterations

The two-by-two intermodality correlation analyses revealed significant results for the atrophy versus hypometabolism analysis only (Fig. 4). Specifically, hypometabolism was correlated to local atrophy in the hippocampus, temporo-parietal cortex, cuneus/precuneus junction, as well as the dorsolateral prefrontal cortex. In contrast, correlations between A β deposition and atrophy or hypometabolism did not reach significance at the $p_{\text{FWE}} < 0.05$ threshold. Using a more permissive statistical threshold of $p_{\text{uncorrected}} < 0.001$, a significant positive correlation was found between A β load and hypometabolism in a small cluster within the retrosplenial cortex/posterior cingulate (MNI coordinates at the peak: $-4, -52, 16$; t value = 4.53; cluster size = 14 voxels).

Regions of interest analyses

The results of the ANOVA comparing the regional W -scores of the three alterations within the regions of interest are displayed in Figure 5. Significant differences were found between the degrees of atrophy, hypometabolism, and A β deposition in all investigated areas, except for the amygdala (Friedman's ANOVA did not reach significance, $p = 0.55$). *Post hoc* comparisons performed with Wilcoxon's signed-rank test showed clearly different gradients according to regions (Fig. 5), with areas of predominant atrophy (such as the hippocampus) and areas of excessive A β deposition compared with atrophy and hypometabolism (such as the prefrontal cortex).

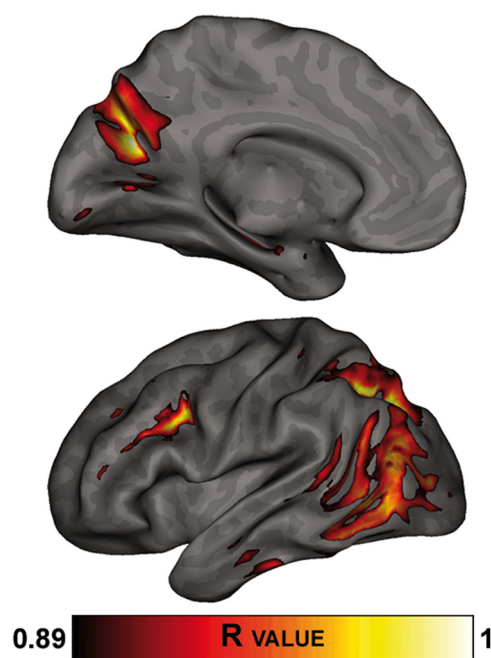


Figure 4. Voxelwise correlations between local degrees of atrophy and hypometabolism in the 20 patients with Alzheimer's disease dementia. The R value of 0.89 used as a threshold in this figure corresponds to the p (FWE corrected) < 0.05 threshold described in Results.

Using the three mean W -scores of each region, Ward's hierarchical clustering method led to the distinction between four subsets of brain areas (Fig. 6, left). A repeated-measures ANOVA was conducted on the mean W -score values from the four sets of regions, with two factors: brain region (with four levels) and imaging modality (with three levels). It revealed a highly significant region \times modality interaction ($F_{(6,152)} = 35.0$; $p < 10^{-5}$), which confirmed the differential hierarchy between the levels of the three markers in these four sets of brain regions (Fig. 6, right). In the hippocampo-amygdala complex, atrophy was marked

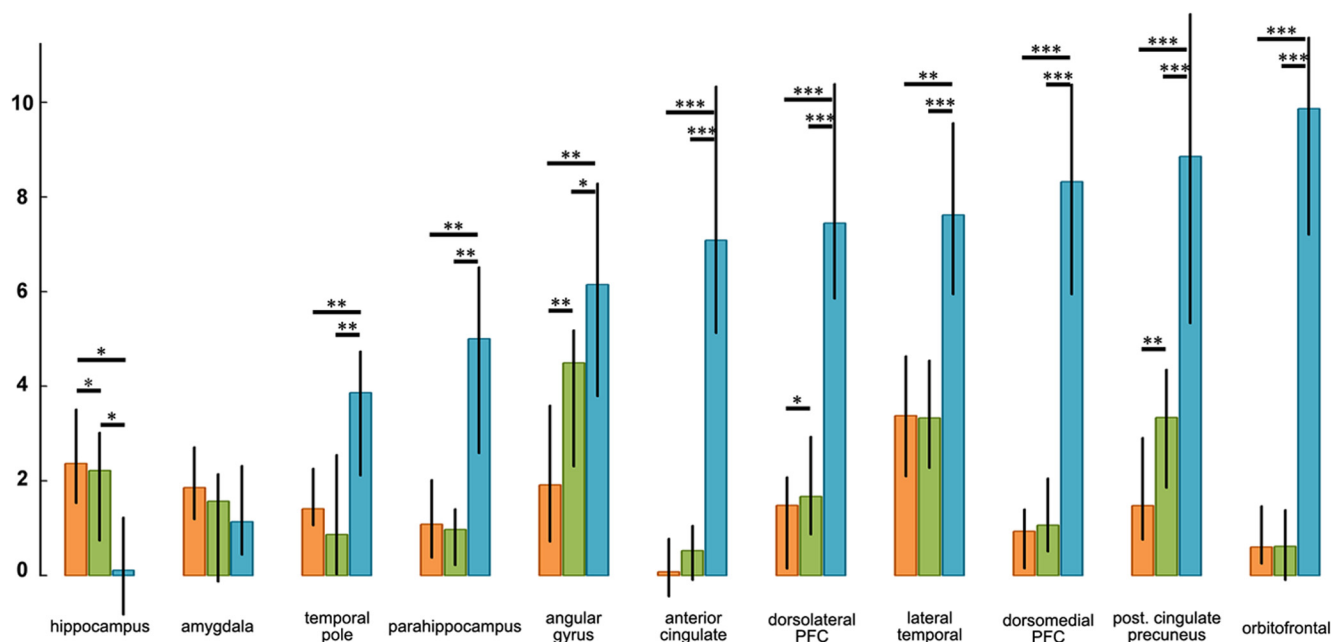


Figure 5. Regions of interest analyses. Local degrees of atrophy (orange), hypometabolism (green), and A β deposition (blue) expressed as mean W -scores were compared using Friedman's ANOVA. When significant ($p < 0.05$), *post hoc* analyses were performed using Wilcoxon's test (* $p < 0.05$; ** $p < 0.005$; *** $p < 0.0001$). Histograms represent median values, and error bars refer to the interquartile range. Regions presented here are ordered by increasing amyloid W -score. post. cingulate, Posterior cingulate cortex; PFC, prefrontal cortex.

(median $W = 2.16$) and significantly exceeded hypometabolism ($W = 1.99$) and A β load ($W = 0.50$). In the parahippocampus and temporal pole, A β load was relatively high ($W = 4.5$) and exceeded local degrees of atrophy ($W = 1.45$) and hypometabolism ($W = 0.78$). In posterior association areas (lateral temporal cortex, angular gyrus, and precuneus/posterior cingulate), all modalities showed high to very high W -scores, with a predominance of A β deposition ($W = 7.36$), exceeding local hypometabolism ($W = 3.69$), itself exceeding atrophy ($W = 2.31$). Last, in frontal areas (dorsolateral prefrontal cortex, dorsomedial prefrontal cortex, orbitofrontal cortex, and anterior cingulate), extreme A β deposition ($W = 7.99$) contrasted to the weakness of structural and metabolic defects (0.61 and 0.85, respectively).

Discussion

In this study, three major brain alterations in Alzheimer's disease, namely gray matter atrophy, hypometabolism, and A β deposition, were measured concomitantly in the same patients with Alzheimer's disease dementia. Their relative expression and relationships were assessed both regionally and voxelwise. We showed marked regional variability in the hierarchy between these different brain alterations with three main profiles: (1) extreme A β deposition with low hypometabolism and atrophy (frontal areas); (2) predominance of A β deposition, together with high hypometabolism and lower but still significant atrophy (posterior association areas); and (3) predominance of atrophy and hypometabolism faced to low A β burden (in medial temporal areas). In addition, local correlations were found between atrophy and hypometabolism within a large parietotemporal network, whereas A β deposition did not correlate to either atrophy or hypometabolism.

One of the most striking results of this study is the considerable discrepancy between A β deposition on the one hand and atrophy and hypometabolism on the other hand in patients with Alzheimer's disease dementia. First, the brain distribution of A β deposition clearly differed from that of the two other alterations

(Fig. 3, middle and right). Second, A β deposition did not correlate to the other alterations. Previous studies assessing the correlations between A β deposition and metabolism in patients with Alzheimer's disease reported conflicting results, and, when significant, correlations were only found in restricted areas, mostly in temporal and parietal cortices (Engler et al., 2006; Edison et al., 2007; Cohen et al., 2009; Forsberg et al., 2010). This is consistent with our finding of a restricted correlation in the posterior cingulate cortex when using a more permissive threshold. As for the link between atrophy and A β deposition, although relationships have been reported in preclinical stages, most studies reported no significant correlations in demented patients (Fagan et al., 2009; Chételat et al., 2010).

These findings are in line with the idea that, at symptomatic stages of the disease, A β does not play a prominent role in ongoing neurodegeneration (Hyman, 2011). Instead, tau pathology, disturbance in axonal transport, or inflammation processes could contribute to brain atrophy and metabolic disruption (see below). They could also be interpreted in the light of current models suggesting a time decay between the appearance of A β plaques, more than one decade before the first symptoms, and hypometabolism and atrophy thought to appear years later, closer to cognitive decline (Jack et al., 2010). Another interesting hypothesis is the possibility of distant, rather than local, effects of A β . Notably, Cohen et al. (2009) observed that metabolism in the precuneus of amyloid-positive patients with Alzheimer's disease was inversely correlated to A β deposition in frontal areas. Bourgeat et al. (2010) also reported a distant relationship (between temporal neocortical A β deposition and hippocampal atrophy), yet the correlation was observed in the cognitively normal group and not in demented patients.

In contrast, atrophy and hypometabolism were found in the same brain areas and were highly correlated (Fig. 4), suggesting that both alterations share at least partly common underlying mechanisms. The direct comparison between both processes also

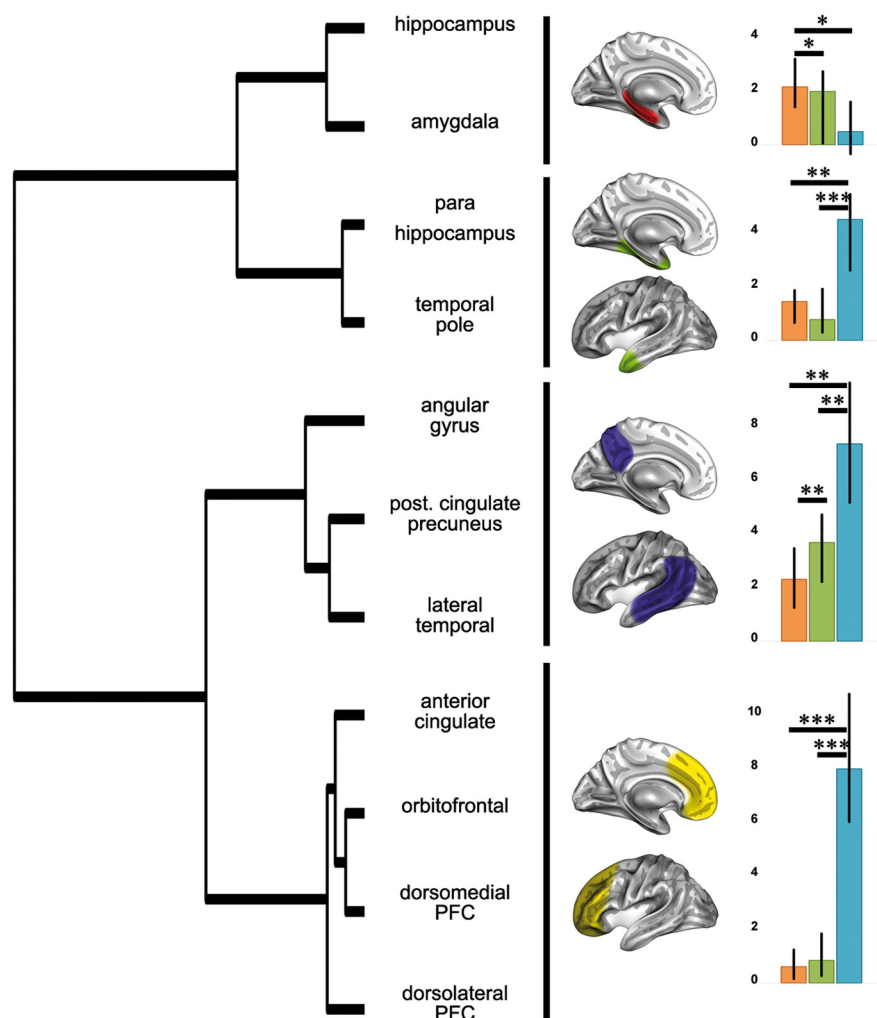


Figure 6. Classification of brain regions according to their degrees of atrophy, hypometabolism, and Aβ deposition. Left, Ward's hierarchical clustering analysis performed on the regions of interest distinguished four subsets of brain areas according to their three mean *W*-scores. Right, For each subset, degrees of atrophy (orange), hypometabolism (green), and Aβ load (blue) were averaged and compared using Friedman's ANOVA and Wilcoxon's test (* $p < 0.05$; ** $p < 0.005$; *** $p < 0.0001$). Histograms represent median values, and error bars refer to the interquartile range. post. cingulate, Posterior cingulate cortex; PFC, prefrontal cortex.

revealed regional differences in their relative degrees (Fig. 3), indicating that additional biochemical mechanisms may amplify or attenuate hypometabolism or atrophy in specific brain regions.

The other notable finding is the regional variability in the hierarchy between the relative degrees of atrophy, hypometabolism, and Aβ deposition, which suggests that these alterations are subtended by multiple region-specific processes. Thus, in contrast to current global models that propose a general chronology in the appearance of the different alterations (independently of brain regions), the present study rather highlights the regional discrepancy in this sequence, suggesting that different subtending processes may be involved in the different brain regions. Although the absolute number of sets of brain regions is arbitrary because it depends on the subjective dendrograms cutoff, our findings revealed clearly distinguishable patterns with three poles: the hippocampus, the posterior association areas, and the frontal cortex.

In the hippocampus, the presence of abnormally phosphorylated tau proteins that aggregate to form neurofibrillary tangles (NFTs) may be one of the processes underlying severe atrophy and moderate hypometabolism despite low Aβ bur-

den. Indeed, neurodegeneration seems to be closely related to tau pathology: progression of both gray matter atrophy (Whitwell et al., 2008) and hypoperfusion (Bradley et al., 2002) through the brain seems to follow NFT spreading as evaluated with Braak staging. In addition, neuronal loss has been shown to correlate with local counts of NFTs in different brain areas, such as the entorhinal cortex (Gómez-Isla et al., 1996) and superior temporal sulcus (Gómez-Isla et al., 1997). Consequently, because NFTs first appear and remain predominant in medial temporal structures as the disease pathology progresses (Braak and Braak, 1991; Delacourte et al., 1999), it is likely that NFTs, rather than Aβ deposition, are responsible for hippocampal atrophy and hypometabolism. However, the possibility of a distant effect of Aβ on the hippocampus cannot be ruled out, especially because the brain structures to which the hippocampus projects the most, i.e., the retrosplenial cortex and medial and orbital prefrontal cortices (Aggleton, 2012), are also those showing the highest Aβ load (Fig. 2, right).

In posterior association areas, several factors have been proposed to induce neuronal dysfunction over and above local Aβ pathology. Notably, recent studies suggested that hypometabolism observed in patients with mild cognitive impairment and Alzheimer's disease dementia (especially in the posterior cingulate cortex) at least partly results from hippocampal atrophy through cingulate bundle disruption (Villain et al., 2008; Choo et al., 2010; Villain et al., 2010). Besides, as mentioned previously, relationships between hypometabolism in the precuneus and prefrontal Aβ load have been observed (Cohen et al., 2009), possibly revealing distant Aβ effects. Actually, the posterior cingulate, precuneus, angular gyrus, and lateral temporal cortex are "cortical hubs" (Buckner et al., 2009), i.e., nodes in brain networks that are tightly connected to numerous brain structures. This may explain their vulnerability to Alzheimer's disease-related pathological processes in widely distributed brain regions.

Last, in frontal areas, Aβ load reached its maxima, whereas atrophy and hypometabolism were minimal. Although already noticed by previous authors (Edison et al., 2007; Jack et al., 2008; Li et al., 2008), the present study provides a quantitative evidence for this discrepancy. This may appear surprising considering studies in animal models showing that severe neurite abnormalities especially develop in the vicinity of Aβ deposits (Tsai et al., 2004) because they are surrounded by a halo of soluble oligomeric forms of Aβ (Koffie et al., 2009), themselves shown to be toxic to synapses (Lacor et al., 2007). Our findings may reflect (1) a lack of deleterious effect of local Aβ deposits in humans, (2) a difference in the timing of the different biomarkers (see above), or (3) the presence of compensation processes. This later hypothesis is supported by pre-

vious functional MRI studies showing greater activations in patients with Alzheimer's disease relative to healthy controls (notably in the frontal cortex), positively associated with cognitive performances and interpreted as the reflect of compensation processes (for review, see Schwindt and Black, 2009). It is thus also possible that, faced with A β deposition, neuronal and synaptic plasticity occurs, allowing to maintain neuronal integrity, glucose consumption, and brain volume.

Note that the patient sample size is relatively limited in the present study compared with large multicenter studies. However, (1) our patients have been thoroughly selected as having a high probability of Alzheimer's disease etiology based on neuroimaging biomarkers as recently recommended, (2) the use of multimodal data obtained in a single laboratory has clear advantage on multicenter data for complex and precise voxelwise analyses as performed here, (3) our findings were confirmed using nonparametric tests, and (4) our results are entirely consistent with previous studies when applicable. Thus, whole-brain patterns of atrophy, hypometabolism, and A β deposition (Fig. 2) are highly similar to those reported in studies from different samples (Buckner et al., 2005; Edison et al., 2007; Alsop et al., 2008; Chételat et al., 2008; Jack et al., 2008; Caroli et al., 2010; Fleisher et al., 2011), and the map of local mismatch between atrophy and hypometabolism (Fig. 3, left) is almost identical to previous studies by Alsop et al. (2008) and Caroli et al. (2010) despite the use of different methodologies and samples. However, we cannot exclude the possibility of a lack of statistical power in the analyses, for instance, to detect subtle relationships between A β deposition and neurodegeneration.

As a whole, our findings highlight the complex local relationships between atrophy, hypometabolism, and A β load and the regional heterogeneity in their hierarchy, suggesting the involvement of different underlying processes. It would be of particular interest to assess these relationships in earlier stages of the disease and to consider not only local but also distant phenomena to get a more comprehensive overview of these different brain alterations and their interactions. Finally, longitudinal studies are warranted to take into account the potential lapse between the different pathological processes while assessing their relationships and eventually propose a more complete, region-specific, model of Alzheimer's disease pathology evolution.

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