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Hippocampal subfield volumetry in Mild Cognitive Impairment, Alzheimer's disease and semantic dementia

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Abstract

Background. Hippocampal atrophy is a well-known feature of Alzheimer's disease (AD), but sensitivity and specificity of hippocampal volumetry are limited. Neuropathological studies have shown that hippocampal subfields are differentially vulnerable to AD; hippocampal subfield volumetry may thus prove to be more accurate than global hippocampal volumetry to detect AD.

Methods. CA1, subiculum and other subfields were manually delineated from 40 healthy controls, 18 AD, 17 amnesic Mild Cognitive Impairment (aMCI), and 8 Semantic Dementia (SD) patients using a previously developed high resolution MRI procedure. Non-parametric group comparisons and receiver operating characteristic (ROC) analyses were conducted. Complementary analyses were conducted to evaluate differences of hemispheric asymmetry and anterior-predominance between AD and SD patients and to distinguish aMCI patients with or without β -amyloid deposition as assessed by Florbetapir-TEP. **Results.** Global hippocampi were atrophied in all three patient groups and volume decreases were maximal in the CA1 subfield (22% loss in aMCI, 27% in both AD and SD; all $p < 0.001$). In aMCI, CA1 volumetry was more accurate than global hippocampal measurement to distinguish patients from controls (areas under the ROC Curve = 0.88 and 0.76, respectively; $p = 0.05$) and preliminary analyses suggest that it was independent from the presence of β -amyloid deposition. In patients with SD, whereas the degree of CA1 and subiculum atrophy was similar to that found in AD patients, hemispheric and anterior-posterior asymmetry was significantly more marked than in AD with greater involvement of the left and anterior hippocampal subfields. **Conclusions.** The findings suggest that CA1 measurement is more sensitive than global hippocampal volumetry to detect structural changes at the pre-dementia stage, although the predominance of CA1 atrophy does not appear to be specific to AD pathophysiological processes.

Keywords: hippocampal subfields, CA1, Mild Cognitive Impairment (MCI), Alzheimer's disease, semantic dementia, Magnetic Resonance Imaging (MRI)

Abbreviations:

A β = β -amyloid;

AD = Alzheimer's Disease;

aMCI = amnesic Mild Cognitive Impairment;

ANOVA = ANalysis Of VAriance;

AUC = Area Under the receiver operating characteristic Curve;

HC = healthy controls;

MRI = Magnetic Resonance Imaging;

NFT = NeuroFibrillary Tangles;

PET = Positron Emission Tomography;

ROC = Receiver Operating Characteristic;

SUV_r = Standardized Uptake Value ratio;

TIV = Total Intracranial Volume.

1. Introduction

Hippocampal atrophy is a major feature of Alzheimer's disease (AD) [1] that strongly correlates with AD neuropathology [2,3]. Using MRI, this atrophy is detectable prior to the diagnosis of AD, at the stage of amnesic Mild Cognitive Impairment (aMCI) [4,5] or even earlier, in asymptomatic elderly up to 10 years before the diagnosis of dementia [6,7]. Hippocampal volume has thus been proposed as a neuroimaging biomarker for early AD diagnosis [8,9]. Yet, the accuracy of this measurement is limited by a moderate sensitivity and a rather low specificity to AD pathophysiological processes [1]. Indeed, hippocampal atrophy has been highlighted in various neurological and psychiatric conditions [10], and notably in other neurodegenerative disorders such as semantic dementia (SD) [11–15].

Interestingly, neuropathological studies have shown a differential vulnerability of hippocampal subfields to AD, with CA1 showing the earliest and strongest changes in terms of both neurofibrillary tangles (NFT) [16,17] and neuronal loss [18,19]. *In vivo* measurement of CA1 atrophy may thus constitute a better surrogate marker for AD pathology than global hippocampal volumetry. Recently MRI acquisition and processing techniques have been developed to assess the hippocampus in more detail, including high-resolution hippocampus scans allowing visualization and measurement of hippocampal subfields [20–22].

In this study, we aimed at identifying the pattern of hippocampal subfield atrophy in patients with AD, aMCI and SD using a previously developed technique based on high-resolution 3-Tesla-MRI and adapted delineation guidelines [23]. In addition we assessed the diagnosis accuracy of these measures, hypothesizing that specific measurement of the most vulnerable subfield(s), e.g. CA1, would be more accurate than global hippocampal volumetry to detect AD-related hippocampal atrophy [20,21].

2. Material and methods

2.1. Participants

Eighty-three right-handed native French-speaking participants from the IMAP project [23] were included in the present study: 40 healthy controls (HC), 17 aMCI patients, 18 AD patients, and eight SD patients (see Table 1). They were all aged over 50 years, had at least 7 years of education and had no history of alcoholism, drug abuse, head trauma or psychiatric disorder.

All patients were recruited from local memory clinics and selected according to corresponding internationally agreed criteria: Petersen's criteria for aMCI [24], NINCDS-ADRDA criteria for probable Alzheimer's disease [25] and Neary et al criteria for SD [26].

HC were recruited from the community and performed in the normal range on a neuropsychological examination assessing multiple domains of cognition including episodic and semantic memory, executive and visuo-spatial functions, language and praxis.

In addition to this clinically based selection, the majority of aMCI and AD patients underwent a Florbetapir PET scan allowing to classify them as β -amyloid ($A\beta$)-positive or $A\beta$ -negative as recommended in the new research criteria for AD [8,9] (see sections 2.3.3 and 2.4 below).

The IMAP Study was approved by regional ethics committee (*Comité de Protection des Personnes Nord-Ouest III*) and is registered with ClinicalTrials.gov (number NCT01638949). All participants gave written informed consent to the study prior to the investigation.

2.2. Neuroimaging data acquisition.

All participants were scanned on the same MRI and PET cameras at the Cyceron center (Caen, France).

2.2.1. MRI data.

A high-resolution proton density-weighted MR sequence, perpendicular to the long axis of the hippocampus (repetition time/echo time = 3500/19 msec; flip angle = 90°; 13 x 2 mm slices with 2

mm gap; in-plane resolution = 0.375 x 0.375 mm) was acquired on a Philips (Eindhoven, The Netherlands) Achieva 3T scanner. A sagittal T1-weighted anatomical image was obtained beforehand using a 3D fast field echo sequence (repetition time/echo time = 20/4.6 msec; flip angle = 20°; 180 x 1 mm slices without gap; in-plane resolution = 1 x 1 mm) for the purpose of PET data preprocessing and Total Intracranial Volume (TIV) measurement (see below).

2.2.2. PET data.

Florbetapir PET scans were acquired on a Discovery RX VCT 64 PET-CT device (General Electric Healthcare) with a resolution of 3.76 x 3.76 x 4.9 mm (field of view = 157 mm). Forty-seven planes were obtained with a voxel size of 2.7 x 2.7 x 3.27 mm. A transmission scan was performed for attenuation correction before the PET acquisition. Participants underwent a 20-min PET scan, starting 50 min after the intravenous injection of ≈ 4 MBq/kg of Florbetapir.

2.3. Neuroimaging data processing.

2.3.1. Hippocampal subfield delineation.

Three regions of interest (subiculum, CA1 and “other” subfields - encompassing CA2-CA3-CA4 and the dentate gyrus) were delineated on both hippocampi of all individual high-resolution MRI images, following guidelines developed in the lab and described in full details elsewhere [23].

These guidelines were based on the anatomical description from an atlas of the human hippocampus [27]. Delineations were performed on slices perpendicular to the long axis of the hippocampus by a single rater, blind to the identity (age, gender, clinical status) of the participants. Illustrations are provided in Figure 1 (see also Supplementary Figure 1 for further details and additional examples).

Unlike other groups with comparable high-resolution anisotropic images [20,21] we delineated subfields along the head and body of the hippocampus (versus in the body only for these previous articles). Yet, because of the difficulty of distinguishing landmarks for subfields delineation in the hippocampal head, we only considered 3 regions of interest (therefore combining CA2-3-4 and the

dentate gyrus in the same “subfield”) to ensure sufficient reliability and reproducibility (versus 4 or 5 subfields in previous studies [20,21]). As previously mentioned [20,22], measurements of hippocampal subfields not only rely on landmarks derived from anatomical atlases, but also on arbitrary rules that are fixed by the investigators to reach a compromise between reliability/reproducibility and validity. We acknowledge that existing protocols show differences in their definition of hippocampal subfields but, in the absence of direct comparison between in vivo imaging methods (ideally including a confrontation to neuropathological gold standard), the importance of these variations and their influence on results remains unknown. However, a recently formed initiative led by experts in the field is specifically meant to address this issue and to potentially develop a unified hippocampal subfield segmentation protocol (to know more about this group, visit <http://www.hippocampalsubfields.com/>).

Moreover, it is to note that high-resolution scans such as those used here are particularly prone to motion artifacts. To prevent the rejection of a large proportion of individuals from analyses, a procedure was settled to obtain high quality data from all individuals participating to imaging studies in our lab (including, but not restricted to the controls and patients from the present article). Indeed, a visual quality check was performed immediately after or within a few days of the MRI acquisition and was carried out by the same person who analyses the data (RLJ). If image quality was considered insufficient for subfield delineation, the scan was repeated within a few weeks during a second MRI session (this occurred for about 15-20% of healthy controls and 30-40% of the patients included in our total imaging cohort). Rarely, a third scan was proposed. In the end, we failed to obtain an image of sufficient quality in some participants (roughly 5-10% of the controls and 15-20% of the patients) as they could not come back for a repeated scan or because the repeated scan(s) was/were still of insufficient quality; in this case, data were disregarded and excluded from all analyses. The 83 participants included in the present study had high quality data.

2.3.2. Total Intracranial Volume (TIV).

Individual TIV values were obtained from the T1-weighted images using the VBM5 toolbox implemented in the Statistical Parametric Mapping software (SPM5; Wellcome Trust Center for Neuroimaging, Institute of Neurology, London, England).

2.3.3. Florbetapir PET.

PET data were processed as described in reference [28]. Briefly, each individual T1-weighted MRI was segmented into gray and white matter using the VBM5 toolbox (<http://dbm.neuro.uni-jena.de/vbm/vbm5-for-spm5/>). These segments were used for partial volume effect correction of raw PET data using the PMOD software. Using the Statistical Parametric Mapping (SPM) software, resulting images were coregistered onto corresponding MRI and normalized into Montreal Neurological Institute (MNI) space using the deformation parameters defined from the VBM procedure. The mean Florbetapir value in the cerebellum GM was extracted for each subject from the normalized TEP images. Each Florbetapir image was then divided by its corresponding mean cerebellar value, resulting in Florbetapir-PET SUVr data. The global neocortical Florbetapir-PET SUVr value was then computed for each subject from the Florbetapir-PET SUVr data using a neocortex mask (including all regions but the cerebellum, hippocampus, amygdala and subcortical grey nuclei).

2.4. Statistical analyses

Raw volumes of each hippocampal subfield and of the whole hippocampus (corresponding to the sum of the three subfields) were first normalized by the TIV to account for inter-individual variability in head size (normalized volume = 1000 x raw volume / TIV) and then transformed into W-scores, i.e. age and gender-adjusted Z-scores [2]. W-scores provide information about the difference between a patient's value and the value that would be expected in the control group for his/her age and gender. In the present study, the use of W-score is of particular interest because of the rather wide age range of the participants and significant age difference between aMCI and both

HC and SD groups (see Table 1). Due to the relative limited size of the patient groups, Kruskal-Wallis non-parametric analyses of variance (ANOVAs) were then performed on hippocampal subfield and whole hippocampus W-scores, and when a significant effect of group was found ($p < 0.05$), Mann-Whitney U tests were used for pairwise comparisons.

As previous studies reported a stronger hemispheric asymmetry and an anterior predominance of hippocampal atrophy in SD compared with AD [11,12], complementary indices were calculated to characterize these gradients within each subfield. First, hemispheric asymmetry was measured as the absolute difference between left and right hippocampal volumes expressed as a percentage of the total volume ($100 \times |\text{right volume} - \text{left volume}| / \text{bilateral hippocampal volume}$). Second, an index of anterior-posterior gradient was calculated as the percentage of each subfield volume located in the anterior hippocampus ($100 \times \text{anterior volume} / \text{total hippocampal volume}$), with the anterior portion corresponding to the hippocampal head [29–31]. These indices were compared between HC, AD and SD using Kruskal-Wallis ANOVA and Mann-Whitney U tests.

Discriminant analyses were performed to assess the ability of each hippocampal subfield and of the whole hippocampus volume to distinguish i) AD patients from HC, ii) aMCI patients from HC, iii) SD patients from AD. Areas Under the receiver operating characteristic Curve (AUC) of global hippocampus versus subfield volumes were then compared to test the hypothesis that subfield are more accurate than global hippocampal volumetry to discriminate between the groups.

Finally, preliminary analyses were conducted to assess the impact of the presence of A β on the different hippocampal volumes. All AD patients ($n=18$) and 15 patients with aMCI (out of 17) underwent a Florbetapir PET scan and were classified using a previously determined Florbetapir neocortical SUVR of 1.1 (see reference [28]). All AD patients were classified as A β -positive and 9 aMCI patients (60 % of aMCI patients who underwent Florbetapir-PET) were classified as A β -positive. A β -positive and A β -negative aMCI did not differ in age, gender or education (all p -values > 0.8). Statistical analyses on hippocampal volumes included both Spearman's correlation

between A β load and hippocampal atrophy within the 15 aMCI patients, and comparisons between A β -positive versus A β -negative aMCI, and between each subgroup and HC. For the latter analysis, because aMCI were significantly older than the whole HC group (see Table 1), a subgroup of age, gender and education-matched HC (n=28) was used.

3. Results

3.1. Atrophy of hippocampal subfields in patients.

Comparisons of W-scores are illustrated in Figure 2. Briefly, the volume of the whole hippocampus was significantly decreased in all three patient groups as compared to HC (mean volume loss = -12% for aMCI, -22% for AD and -17% for SD). In aMCI, atrophy was highly significant in CA1 (-22%, $p < 0.001$) and to a lesser extent in the subiculum (-17%, $p = 0.01$). In AD, volume decreases were highly significant ($p < 0.001$) for all three subfields, but atrophy in the other region (-17%) was significantly lower ($p < 0.005$ using Wilcoxon rank-sum test) than in both CA1 and subiculum (both -27%). In the SD group, both CA1 and subiculum volumes were significantly reduced relative to HC (-27% and -24% respectively, both $p < 0.001$). Similarly to the findings in the AD group, comparing degrees of atrophy (expressed as W-scores) between the three subfields in SD showed that both CA1 and subiculum were significantly more affected than the other subfield (both $p < 0.05$ Wilcoxon rank-sum test). None of the volumes differed between AD and SD patients (all p values > 0.15). Note that the main results were unchanged when performing comparisons on raw volumes or TIV-normalized volumes instead of age- and gender-adjusted W-scores as presented here (see supplementary figure 2).

3.2. Hemispheric asymmetry and anterior-posterior gradient in AD and SD.

The results of the between-group comparisons of the hippocampal indices are reported in Table 2. Greater hemispheric asymmetry was found in SD patients compared to HC for all volumes (all p

values < 0.03). Asymmetry was significantly stronger in SD than in AD for the whole hippocampus ($p=0.001$), subiculum ($p=0.02$), other ($p=0.01$) and a trend was found for CA1 ($p=0.06$).

Finally, the anterior-posterior index was significantly lower in SD compared to both HC and AD for the whole hippocampus, CA1 and subiculum, indicating a predominance of atrophy in the anterior hippocampus in SD.

3.3. Evaluation of diagnostic accuracy

For the AD versus HC discrimination, the AUC of all subfields was significantly higher than 0.5 (mean [95% CI]=0.92 [0.85-0.99] for CA1, 0.88 [0.76-1] for subiculum, 0.81 [0.67-0.94] for other) but none of them was significantly higher than that of the whole hippocampus (0.91 [0.78-1]). For the discrimination between HC and aMCI, the AUC of CA1 (0.88 [0.78-0.98]) was significantly higher than 0.5 and higher than the AUC of the whole hippocampus (0.76 [0.60-0.93], $p=0.05$; see Figure 3) whereas the subiculum AUC (0.74 [0.59-0.90]) was significantly higher than 0.5 but did not perform better than the whole hippocampus, and the AUC for the other subfields was not significantly different from 0.5 (0.53 [0.34-0.72]). Finally, none of the volume measurements allowed to separate AD from SD as none of the AUCs was significantly different from 0.5: 0.51 [0.27-0.74] for CA1, 0.64 [0.43-0.85] for subiculum, 0.69 [0.45-0.94] for other, and 0.69 [0.47-0.90] for the whole hippocampus.

3.4. Volume of hippocampal subfields versus neocortical amyloid load in aMCI patients.

A Kruskal-Wallis ANOVA with three groups (HC, A β -positive aMCI, A β -negative aMCI) was performed on TIV-normalized hippocampal measurements. A significant effect of group was found for CA1 ($H=13.8$; $p=0.001$), subiculum ($H=5.99$; $p=0.05$) and global hippocampus ($H=5.98$; $p=0.05$). Post-hoc analyses revealed that A β -positive aMCI had significantly lower CA1 ($p<0.001$), subiculum ($p=0.01$) and hippocampal volumes ($p=0.01$) as compared to HC, while A β -negative aMCI only showed lower CA1 volume ($p=0.04$) as compared to HC, as illustrated in Figure 4. The direct comparison between A β -positive aMCI and A β -negative aMCI did not show any significant

difference (all p values >0.25). Similarly, none of the volumes significantly correlated with Florbetapir neocortical SUVR in the whole group of 15 aMCI (CA1: Spearman's $\rho = -0.34$, $p = 0.22$; subiculum $\rho = -0.26$, $p = 0.34$; other $\rho = -0.01$, $p = 0.96$; whole hippocampus $\rho = -0.23$, $p = 0.41$). All the results remained unchanged when using W-scores instead of TIV-normalized volumes (data not shown).

4. Discussion

Using a high-resolution sequence on a 3T MRI scanner, hippocampal subfields were delineated in healthy controls and patients with aMCI, AD or SD. Analyses showed that the pathologies had a differential effect on the hippocampal subfields, with a preferential involvement (ie. stronger and earlier atrophy) of CA1 and subiculum. This pattern was not discriminant at the dementia stage as i) no differences were found between AD and SD and ii) hippocampal subfields did not perform better than the whole hippocampus in discriminating AD from HC. By contrast, at the aMCI stage, CA1 volume loss predominated and was found to better discriminate aMCI patients from HC than global hippocampal volume.

4.1. Hippocampal subfield atrophy in AD

The finding of a differential atrophy of the hippocampal subfields in AD, with a stronger and earlier alteration of CA1 and subiculum, is in agreement with neuropathological studies [16–19]. For instance, West et al [18] reported a reduction in the number of neurons in AD patients compared to HC of 68% in CA1, 47% in the subiculum and 25% in the hilus of the dentate gyrus. The higher vulnerability of CA1 neurons is likely to be at least partly due to the neurofibrillary pathology as the CA1 subfield is the first hippocampal area to be affected by NFT [16,17]. Moreover, several studies have reported a strong negative correlation between neuronal counts and NFT number in the CA1 subfield [32,33] suggesting that the progression of brain atrophy and NFT are strongly associated [34].

Using a variety of neuroimaging approaches such as radial atrophy [35], large-deformation high-dimensional brain mapping [36], voxel-based morphometry [37] or manual delineation [20], the predominance of atrophy in CA1 (and to a lesser extent in the subiculum) in patients with clinical AD or aMCI has already been shown *in vivo*. In the present study, we confirmed this point using a refined methodology in patients with both a clinical diagnosis of AD and a positive Florbetapir PET-scan. Our study showed that all subfields were atrophied at this AD dementia stage and subfield measurements were not more accurate than the global hippocampus in discriminating AD from HC. By contrast, and in agreement with previous volumetric studies [20,21,38], we showed that subfield measurements were more accurate than global hippocampal volumetry to differentiate aMCI from HC, highlighting the interest of these methods for early AD detection. However, discrepancies exist as regards to the area of largest difference between HC and aMCI patients (CA1 [21], CA1-CA2 transition area [20] or subiculum [38]), probably reflecting variations in the anatomical landmarks used for subfield delineation as further discussed in previous publications [22,23] (see the method section 2.3.1 for further discussion).

4.2. Comparison of the pattern of hippocampal atrophy in AD versus SD

AD and SD patients did not differ in terms of subfield volumetry. Although the absence of significant difference could be due to a lack of statistical power, preferential atrophy of both CA1 and subiculum were found in both groups and this finding is consistent with a recent surface-based study of the hippocampus in SD [39]. Yet, the pathological substrate of hippocampal atrophy in SD is not clear, notably because patients with SD can present with heterogeneous pathological features [40]. Despite this variability, a severe neuronal loss in the CA1 subfield together with a preservation of neurons in the dentate gyrus, consistent with our imaging findings, was reported independently of the histopathological subtypes [40].

Contrastingly, hippocampal atrophy differed between SD and AD patients in terms of hemispheric and anterior-posterior asymmetry, in agreement with previous reports [11,12,14,15], and we

showed that these two gradients were mostly independent of subfields. Previous authors [11,14,15] hypothesized that this asymmetric nature of hippocampal atrophy in SD could partly explain the intriguing relative preservation of episodic memory in SD as compared to AD [41] in spite of similarly severe hippocampal atrophy [42]. According to these authors, the relatively spared hippocampal areas (posterior areas and most of the time the right hippocampus) in SD could therefore be sufficient to support essential episodic memory abilities [11,14,15]. Alternatively, it has been proposed [12] that differential alterations of cortical areas that are crucial for episodic memory, such as the posterior cingulate cortex (PCC) that shows a strong hypometabolism in AD but not in SD, are more likely to account for the differential alteration of episodic memory between AD and SD. Yet, these two hypothesis are not necessarily independent as hippocampal atrophy has been shown to induce PCC hypometabolism through a disconnection process in AD [43]. Besides, the connectivity of the hippocampus varies along its anterior-posterior axis, with the posterior part being more strongly connected to the PCC [44]. It is thus possible that the predominance of atrophy in the anterior hippocampus in SD at least partly accounts for differences in distant cortical alterations (e.g. the lack of PCC hypometabolism) compared to AD, themselves underlying differences in the cognitive deficits. This would also be consistent with the idea that SD and AD target two large-scale brain networks that would underlie different cognitive functions and differentially include anterior versus posterior hippocampi [45].

4.3. Relationships between hippocampal subfield atrophy and A β in aMCI

Because the presence of A β deposition significantly increases the likelihood of having AD pathology in patients with aMCI according to current diagnosis recommendations [8,9], we also investigated hippocampal subfield atrophy in aMCI patients as a function of A β load. We did not find a significant influence of the presence of A β on the pattern of hippocampal subfield atrophy as both A β -positive and A β -negative aMCI showed a predominant atrophy of the CA1 subfield (Figure 4). However, these findings should be considered as preliminary given the small size of the

aMCI samples when dichotomized in A β -positive and A β -negative. Yet, this finding is in line with recent studies showing that AD-type atrophy and/or hypometabolism can be found in both A β -negative healthy controls [46] and A β -negative MCI patients [47]. While it may reflect methodological issues related to biomarker measurements [48], it is possible that CA1 atrophy in A β -negative patients reflect non-AD pathophysiological processes. This would be consistent with the finding of CA1 atrophy in other disorders [2,49,50] and the idea that aMCI in A β -negative patients is unlikely due to AD etiology [9]. Alternatively, recent findings suggest that neuronal injury biomarkers known to be closely related to tau pathology such as hippocampal atrophy (see above), may occur independently and possibly prior to A β in the course of AD [51,52]. This would rather support hypotheses suggesting that tau pathology could occur independently from A β accumulation [53], than the amyloid hypothesis where tau-related neurodegeneration is supposed to appear downstream to A β . Further investigations are needed to test these hypotheses but also to replicate our findings regarding the relationships between hippocampal subfield atrophy and A β in a larger sample given the limited size of our aMCI subgroups.

It should also be noted that subfield volumetry as performed in the present study only provides approximations of the exact volumes of the hippocampal subfields, based on anatomical landmarks derived from histological atlases (see the method section 2.3.1 for further discussion).

4.4. Conclusion

Overall, our findings in AD and SD are consistent with the topography of neuronal loss described in post-mortem studies as well as with previous imaging studies that used different methods. They suggest that hippocampal subfield volumetry is a promising biomarker for early AD detection at a prodementia or even presymptomatic stage, especially with the widespread use of high-resolution MR sequences in the last years [20,22] and the on-going development of automatic subfield segmentation procedures [54,55]. Further studies are therefore needed to assess the diagnostic and

prognostic accuracies of this novel technique in larger samples and in comparison to other established AD biomarkers.

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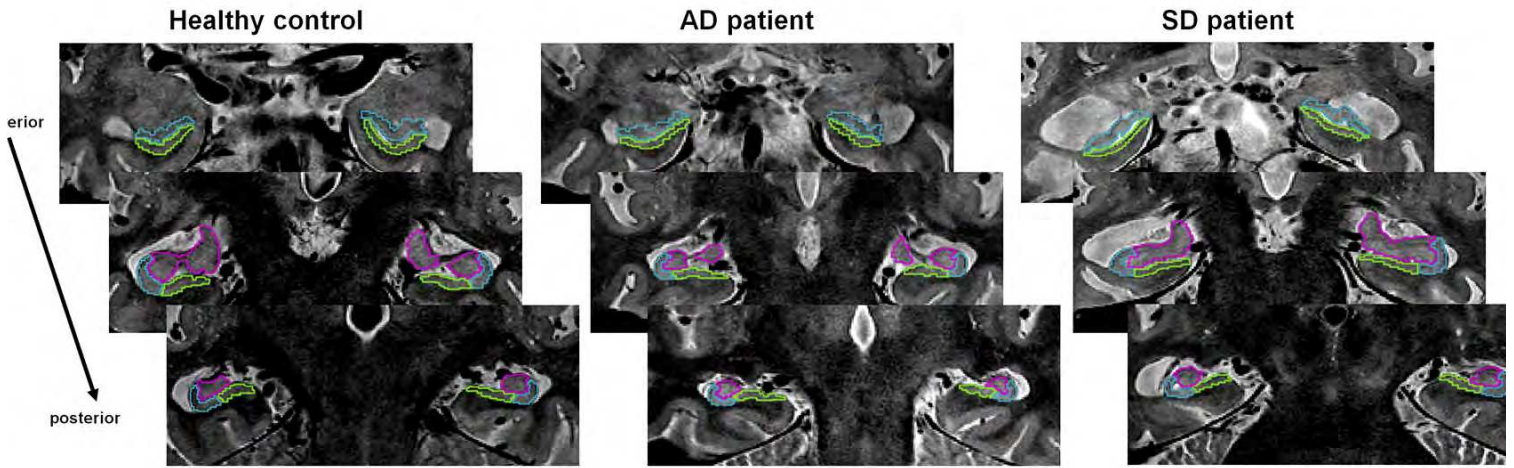
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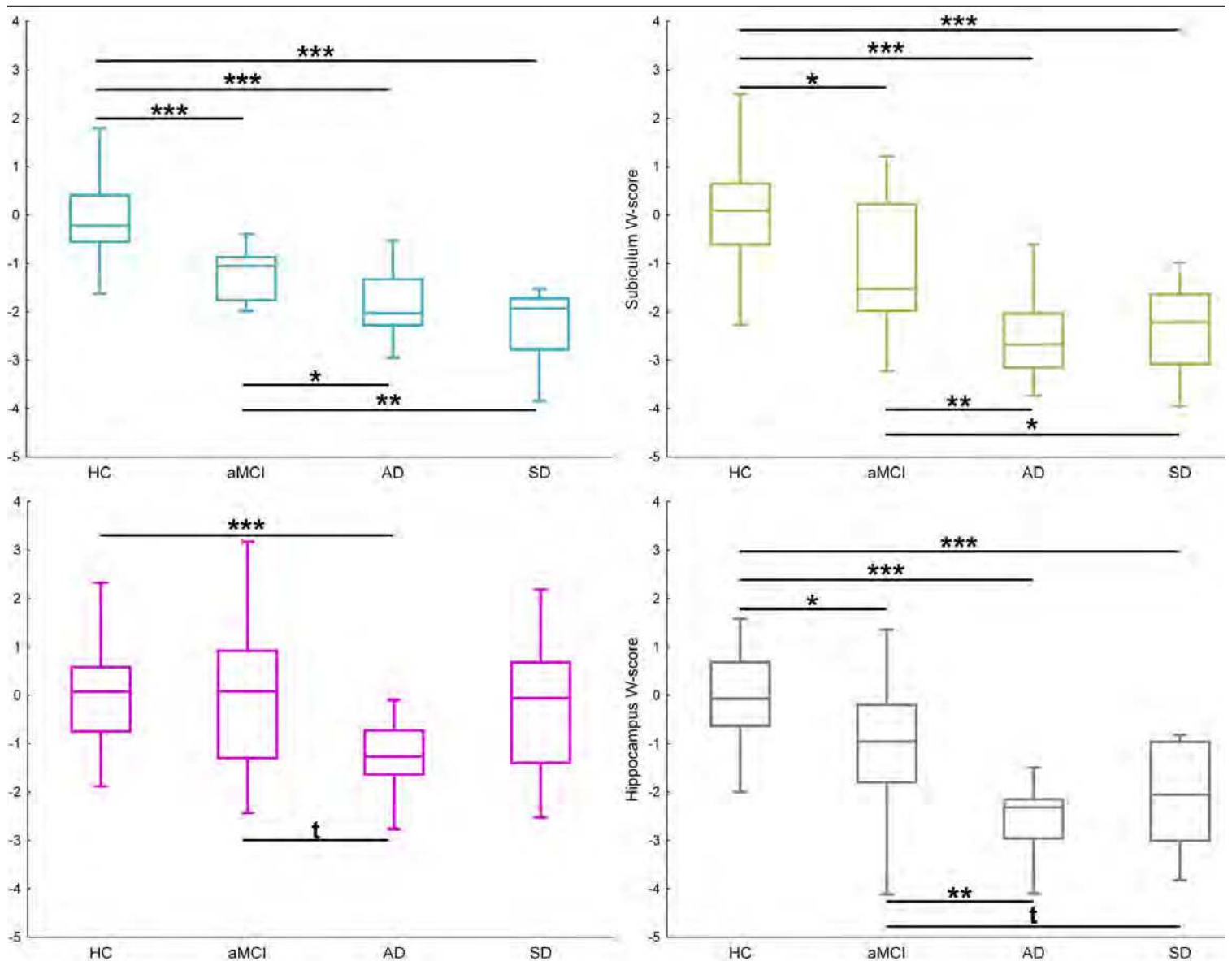
FIGURES

Figure 1. Illustration of hippocampal subfield delineation.



Three regions were manually delineated within each hippocampus: CA1 (blue), subiculum (green) and other (pink). Subfields were delineated on 9 slices on average; for the purpose of illustration, examples are displayed on three slices along the anterior-posterior axis of the hippocampus. Images correspond to a healthy control (left), a patient with Alzheimer's disease (middle) and a patient with semantic dementia (right). Images are in the neurological convention (right is right). Additional examples are available in supplementary figure 1.

Figure 2. Between-group comparisons of hippocampal measurements.

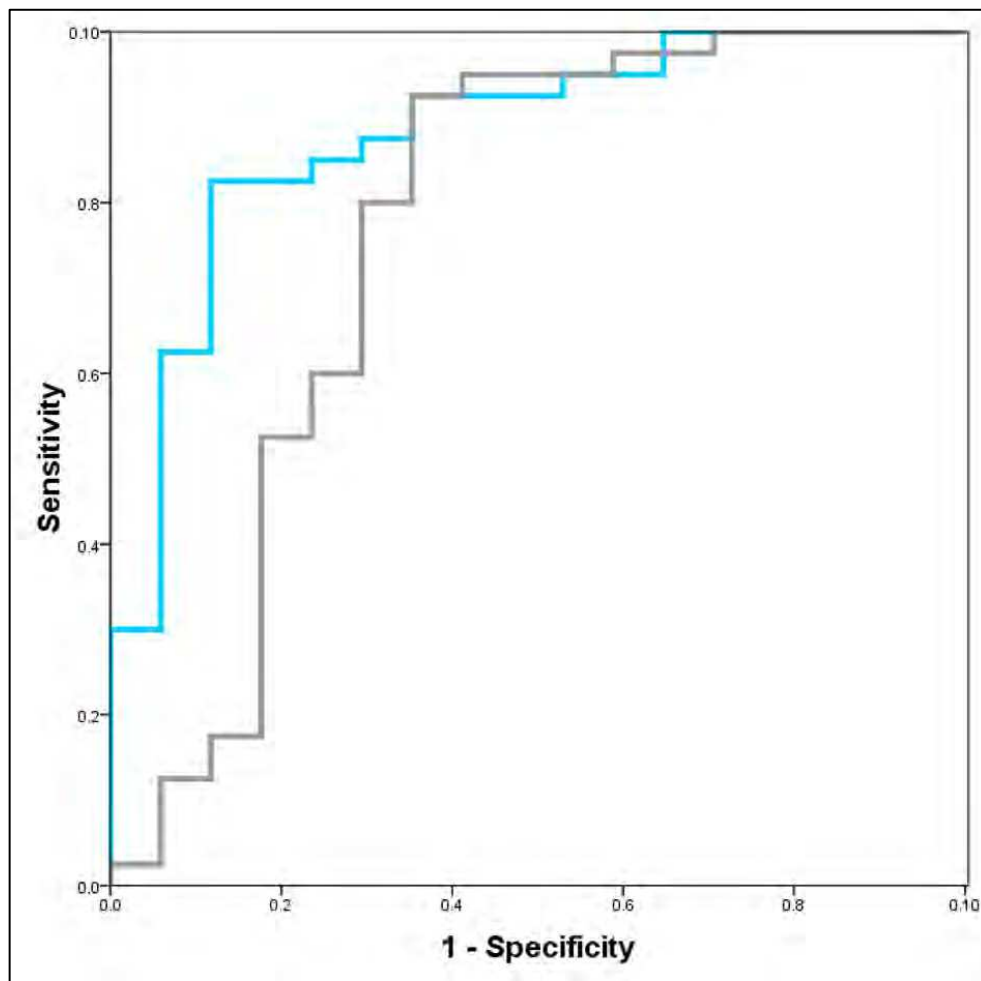


Abbreviations: HC = healthy controls; aMCI = amnesic mild cognitive impairment; AD = Alzheimer's disease; SD = semantic dementia.

Volumes are expressed as W-scores (ie age- and gender-adjusted Z-scores as compared to the control group).

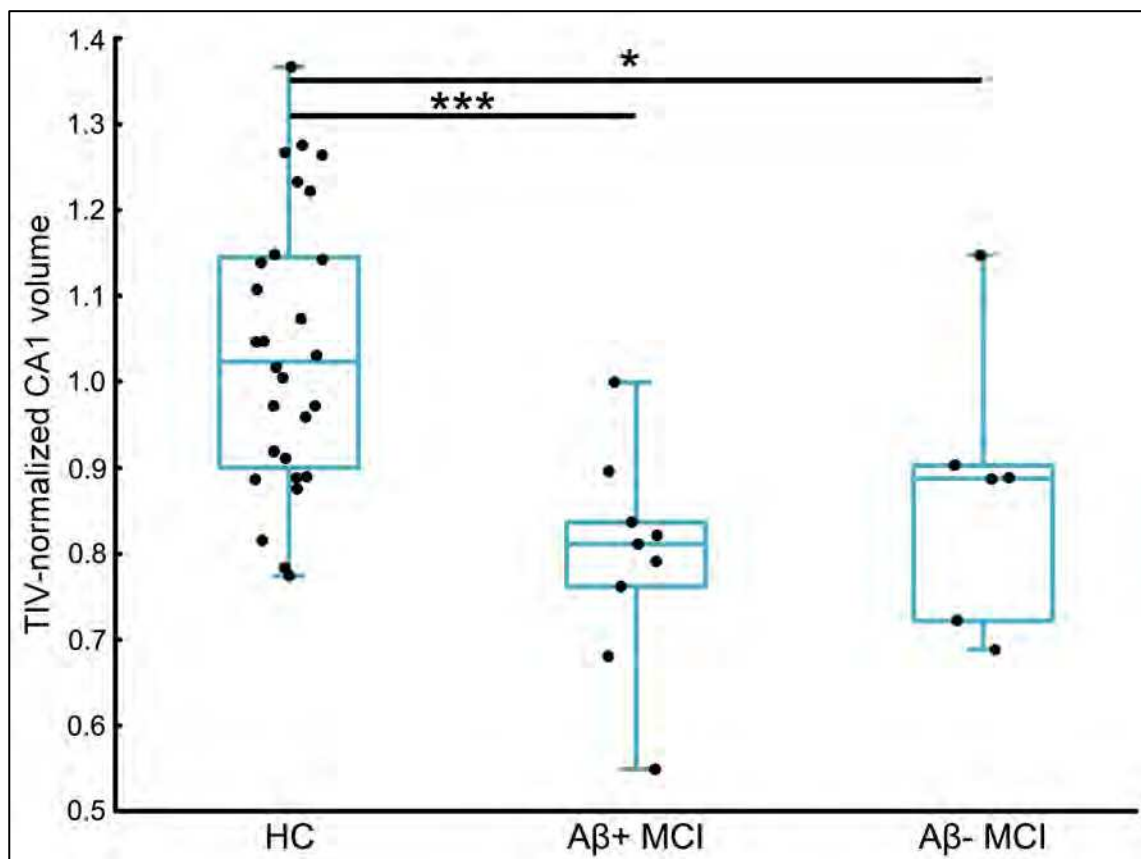
Kruskall-Wallis ANOVA was significant for all regions and post-hoc tests were performed with the Mann-Whitney test (t: $p < 0.10$; *: $p < 0.05$; **: $p < 0.01$; ***: $p < 0.001$).

Figure 3. ROC Curve for CA1 and total hippocampal volume in aMCI patients versus HC.



ROC curves for CA1 (blue) and total hippocampal volume (gray) in the comparison between healthy controls (HC) and patients with amnesic Mild Cognitive Impairment (aMCI). The area under the curve (AUC) is significantly higher ($p = 0.05$) for CA1 (AUC = 0.881) than for total hippocampal volume (0.763).

Figure 4. CA1 atrophy in aMCI patients classified as A β -positive or A β -negative according to their neocortical Florbetapir SUVr.



Abbreviations: HC = healthy controls; A β + aMCI = β amyloid-positive patients with amnesic mild cognitive impairment; A β - aMCI = β amyloid-negative patients with amnesic mild cognitive impairment

For this analysis, a subsample of 28 age-, gender-, and education-matched HC was selected.

Kruskall-Wallis ANOVA was significant ($p=0.001$) and post-hoc tests were performed with the Mann-Whitney test (*: $p<0.05$; ***: $p<0.001$).

TABLES

Table 1. Demographic characteristics of participants.

	HC (n=40)	aMCI (n=17)	AD (n=18)	SD (n=8)	Group comparison
Age	66 (56, 72)	72 (69, 75)	66 (58, 76)	62.5 (59.5, 65.5)	$p_{ANOVA}=0.027$
Median (IQR)					aMCI>HC; aMCI>SD
Gender	23 (58%)	9 (53%)	12 (67%)	5 (63%)	$p_{Chi-squared}=0.86$
Females: n (%)					
Years of education	12 (9.5, 15)	10 (8, 12)	10 (7, 14)	12 (5.8, 15.5)	$p_{ANOVA}=0.19$
Median (IQR)					
MMSE	30 (29, 30)	27 (26, 28)	20.5 (20, 24)	-	$p_{ANOVA}<0.001$
Median (IQR)					HC>aMCI>AD
Mattis	143 (142, 144)	137 (134, 138)	123 (114, 128)	118 (111, 125.5)	$p_{ANOVA}<0.001$
Median (IQR)					HC>aMCI>AD, SD ^a

Abbreviations: HC = healthy controls; aMCI = amnesic mild cognitive impairment; AD = Alzheimer's disease; SD = semantic dementia; IQR = interquartile range; MMSE = mini-mental state examination; ANOVA = Analysis of variance.

All variables were compared using non-parametric tests.

a: Mattis scores were lower in AD and SD groups as compared to HC and aMCI but no difference was found between AD and SD.

Table 2. Hemispheric asymmetry and anterior-posterior gradients of subfield atrophy in AD and SD patients.

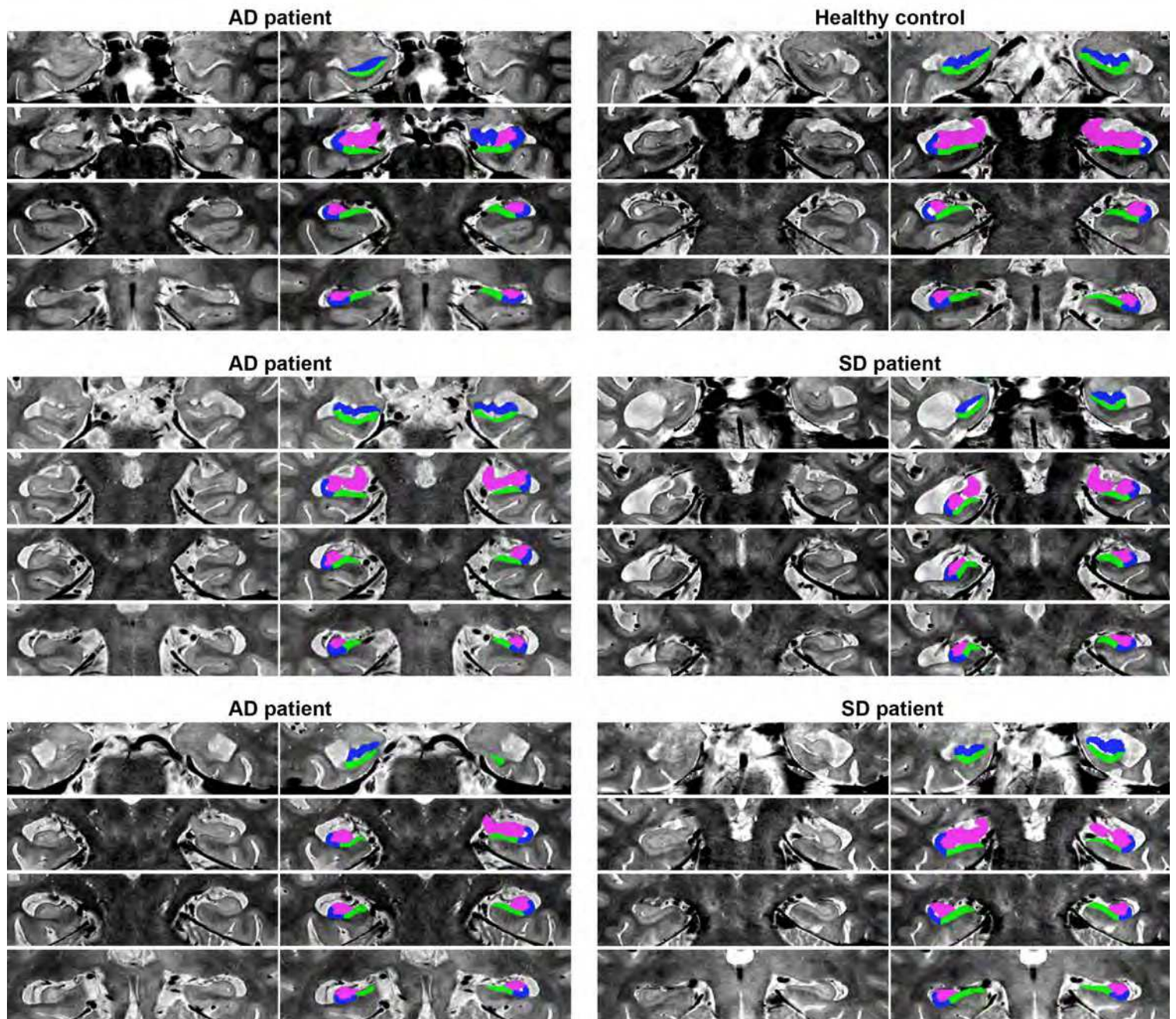
	Values: median (IQR)			ANOVA	Statistical comparisons		
	HC	AD	SD		HC / AD	HC / SD	AD / SD
Hemispheric asymmetry							
CA1	5.4 (1.8, 8.5)	7.8 (4.2, 17.0)	15.3 (13.1, 18.5)	H = 14.7 p < 10⁻³	Z = 1.94 p = 0.05	Z = 3.67 p < 10⁻³	Z = 1.86 p = 0.06
Subiculum	3.8 (1.6, 6.5)	6.6 (1.8, 8.1)	17.7 (12.7, 23.8)	H = 16.1 p < 10⁻³	Z = 1.64 p = 0.10	Z = 3.91 p < 10⁻³	Z = 2.42 p = 0.02
Other	5.2 (2.8, 9.8)	5.1 (1.9, 8.1)	11.4 (7.2, 14.4)	H = 7.2 p = 0.03	Z = -0.82 p = 0.42	Z = 2.31 p = 0.021	Z = 2.53 p = 0.01
Global hippocampus	2.1 (1.2, 4.0)	4.7 (2.8, 7.1)	14.2 (11.2, 16.1)	H = 27.1 p < 10⁻³	Z = 3.17 p = 0.002	Z = 4.41 p < 10⁻³	Z = 3.25 p = 0.001
Anterior-posterior gradient							
CA1	66.2 (63.4, 70.0)	66.6 (59.7, 70.1)	56.5 (52.9, 57.6)	H = 14.7 p < 10⁻³	Z = -0.33 p = 0.74	Z = -3.80 p < 10⁻³	Z = 3.08 p = 0.002
Subiculum	57.2 (53.2, 61.5)	60.1 (53.3, 65.6)	52.4 (44.8, 54.9)	H = 8.3 p = 0.02	Z = 1.02 p = 0.31	Z = 2.59 p = 0.01	Z = 2.47 p = 0.01
Other	62.8 (57.5, 66.7)	65.8 (57.1, 70.4)	59.7 (55.5, 61.9)	H = 3.03 p = 0.22	-	-	-
Global hippocampus	62.4 (59.0, 65.1)	64.6 (58.2, 68.5)	55.6 (52.0, 59.0)	H = 9.3 p = 0.009	Z = 0.60 p = 0.55	Z = 2.95 p = 0.003	Z = 2.53 p = 0.01

Abbreviations: HC = healthy controls; AD = Alzheimer's disease; SD = semantic dementia; IQR = interquartile range. ANOVA were conducted with the Kruskal & Wallis H test and when significant ($p < 0.05$), two-by-two comparisons were assessed with the Mann-Whitney test. Significant ($p < 0.05$) results are shown in bold.

Increased hemispheric asymmetry indicates a stronger left / right volume difference but not the direction of this difference. Increased anterior-posterior gradient index indicates an increase in the volume of the anterior hippocampal compared to that of the total hippocampal volume. See main text (section 2.4) for further information.

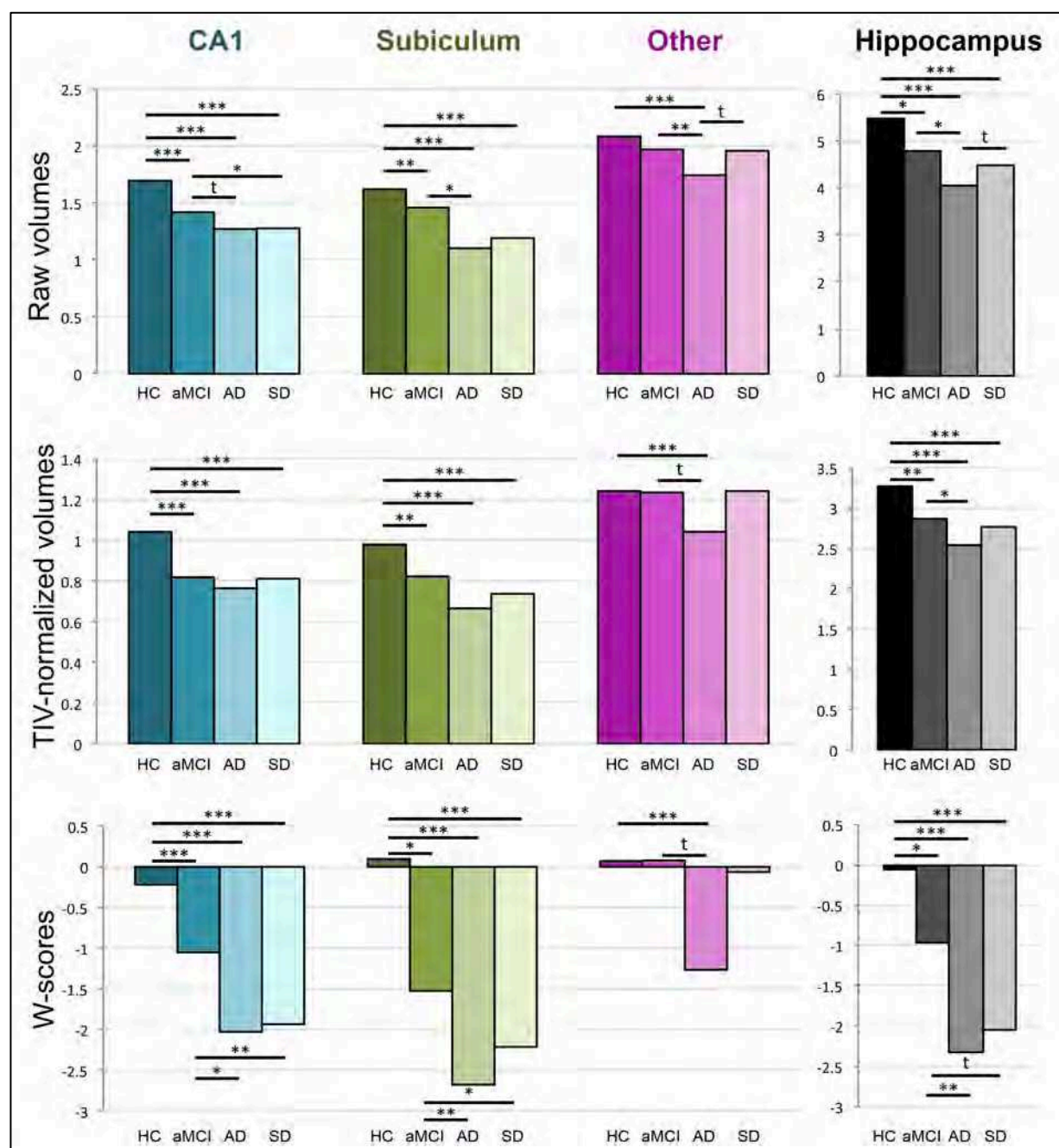
SUPPLEMENTARY MATERIALS

Supplementary figure 1



Additional examples of hippocampal subfield delineation in 3 patients with Alzheimer's disease (AD), two patients with semantic dementia (SD) and a healthy control. For each individual, different slices are shown with and without annotations to enable a better visualization of the hippocampus.

Supplementary figure 2. Influence of the correction for total intracranial volume and demographic factors on group comparisons.



Abbreviations: HC = healthy controls; aMCI = amnesic mild cognitive impairment; AD = Alzheimer's disease; SD = semantic dementia; TIV = total intracranial volume.

In the upper row, raw volumes were directly compared between groups. On the middle row, volumes were first normalized by the TIV using the following formula: $\text{normalized volume} = 1000 \times \text{raw volume} / \text{TIV}$. On the lower row, TIV-normalized volumes were transformed into W-scores (i.e. age and gender-adjusted Z-scores). This last row shows the same results as those presented in the main manuscript (figure 2) but are presented here for the sake of comparison with the other conditions.

Pairwise comparisons were performed with the Mann-Whitney test (t: $p < 0.10$; *: $p < 0.05$; **: $p < 0.01$; ***: $p < 0.001$). Bars represent the median value for each group.