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LONGITUDINAL BRAIN METABOLIC CHANGES FROM AMNESTIC MILD COGNITIVE IMPAIRMENT TO ALZHEIMER'S DISEASE

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ABSTRACT

A sensitive marker for monitoring progression of early Alzheimer's Disease (AD) would help to develop and test new therapeutic strategies. The present study aimed at investigating brain metabolism changes over time, as potential monitoring marker, in patients with amnesic Mild Cognitive Impairment (aMCI), according to their clinical outcome (converters or non-converters), and in relation to their cognitive decline. Seventeen aMCI patients underwent MRI and ^{18}F FDG-PET scans both at inclusion and 18 months later. Baseline and follow-up PET data were corrected for partial volume effects and spatially normalized using MRI data, scaled to the vermis and compared using SPM2. 'PET-PAC' maps reflecting metabolic percent annual changes were created for correlation analyses with cognitive decline. In the whole sample, the greatest metabolic decrease concerned the posterior cingulate-precuneus area. Converters had significantly greater metabolic decrease than non-converters in two ventro-medial prefrontal areas, the subgenual (BA25) and anterior cingulate (BA24/32). PET-PAC in BA25 and BA24/32 combined allowed complete between-group discrimination. BA25 PET-PAC significantly correlated with both cognitive decline and PET-PAC in the hippocampal region and temporal pole, while BA24/32 PET-PAC correlated with posterior cingulate PET-PAC. Finally, the metabolic change in BA8/9/10 was inversely related to that in BA25 and showed relative increase with cognitive decline, suggesting that compensatory processes may occur in this dorso-medial prefrontal region. The observed ventro-medial prefrontal disruption is likely to reflect disconnection from the hippocampus, both indirectly through the cingulum bundle and posterior cingulate cortex for BA24/32, and directly through the uncinate fasciculus for BA25. Altogether, our findings emphasize the potential of ^{18}F FDG-PET for monitoring early AD progression.

Keywords:

amnesic Mild Cognitive Impairment, ¹⁸F-DG-PET monitoring, ventro-medial prefrontal cortex, longitudinal study

INTRODUCTION

To develop and test new therapeutic strategies for Alzheimer's Disease (AD), sensitive markers for monitoring disease progression are urgently needed especially at its earliest stages, when neuropathological damage is still confined.

Positron Emission Tomography with 2-[¹⁸F]-Fluoro-2-Deoxy-D-Glucose (¹⁸FDG-PET) is exquisitely sensitive to early AD-related brain changes. Significant hypometabolism can be detected in patients with amnesic Mild Cognitive Impairment (aMCI) that best represents the pre-dementia stage of AD Petersen, 2005. The earliest metabolic changes involve the precuneus - posterior cingulate cortex (PCC) and temporo-parietal areas (Salmon *et al.*, 1994; Minoshima *et al.*, 1997; Drzezga *et al.*, 2003; Nestor *et al.*, 2003a; Chételat *et al.*, 2003b; Nestor *et al.*, 2003b; Nestor *et al.*, 2004; Mosconi, 2005; Ishii *et al.*, 2005; Kawachi *et al.*, 2006). By contrast, the frontal cortex appears involved at the dementia stage (Minoshima *et al.*, 1997; Desgranges *et al.*, 1998; Alexander *et al.*, 2002; Herholz *et al.*, 2002), suggesting that metabolic changes should be detectable from aMCI to clinically probable AD.

Nonetheless, little is known about the accuracy of ¹⁸FDG-PET to monitor the progression of early AD and only one previous longitudinal ¹⁸FDG-PET study in aMCI patients has been published so far (Drzezga *et al.*, 2003). This study highlighted significantly greater metabolic decreases in the right middle frontal gyrus in those patients who converted to clinically probable AD as compared to non-converters over a one-year follow-up period. However, whilst clinically meaningful, the dichotomous approach used in this study (i.e. comparing rapid converters to non-converters) is limited by the fact that non-converters do include patients who will later progress to AD. A comprehensive approach should also consider all aMCI patients, whether they rapidly convert or not, assessing in a whole sample analysis brain metabolic changes

in relation to cognitive decline over time (Chételat *et al.*, 2005a). Indeed, these two approaches would appear as complementary as the former would allow identifying specific changes occurring in aMCI patients while they convert to clinically probable AD while the latter would inform on changes characterizing rapid cognitive decline whatever the clinical outcome at the end of an arbitrary-defined follow-up period.

Our main objective with the present study was therefore to further investigate the brain pattern of metabolic changes over the course of early AD using these two complementary approaches, i.e. comparing converters to non-converters in a standard fashion, but also across the whole sample in relation to cognitive decline, implementing methodology specially designed for this purpose. Furthermore, thanks to supplementary analyses, we aimed at assessing their discriminant accuracy for monitoring the progression to early AD and investigating the mechanisms underlying these metabolic changes.

METHODS

Patients

The present sample partly overlaps with those of our previous publications using baseline PET data (Chételat *et al.*, 2003a; Chételat *et al.*, 2003b; Chételat *et al.*, 2005a; Mevel *et al.*, 2007) or longitudinal magnetic resonance imaging (MRI) data (Chételat *et al.*, 2005b; Chételat *et al.*, 2008), although only those patients with both baseline and follow-up MRI and PET data were included in this study. Briefly, the 17 aMCI patients included here were all recruited through a memory clinic, and all complained of memory impairment. They were right-handed, aged over 55 years and had at least 7 years of education (see Table 1 for demographic and clinical characteristics). They underwent medical, neurological, neuropsychological, and neuroradiological examinations, and were selected according to current criteria of aMCI, i.e. isolated episodic memory deficits (<1.5 SD of the normal mean for age and education), normal performances in other areas of cognition and in global cognition (assessed with MMSE and Mattis scales), and NINCDS-ADRDA criteria for probable AD (McKhann *et al.*, 1984) not met (see Chételat *et al.*, 2005a for details). According to the Declaration of Helsinki, each patient gave written informed consent to participate in the study, which was approved by the regional ethics committee.

Using the same neuropsychological battery as used at inclusion, all aMCI patients were evaluated every 6 months over an 18-month follow-up period to assess whether they met NINCDS-ADRDA criteria of probable AD or not; at the end of the follow-up period, patients were classified as converters or non-converters, respectively. Patients were declared as converters if they had impaired performances (more than 1.5 SD below the normal means according to age and education when

available) in at least one of general intellectual function scales as well as in at least two areas of cognition including memory, leading to impaired daily activities as judged by the clinicians from the consultation interviews. Moreover, as an index of cognitive decline, a Mattis - Percent Annual Change (Mattis-PAC) was obtained for each patient. This index was calculated by first modelling a simple linear regression from Mattis scores collected at each neuropsychological evaluation ($y=ax+b$; where y =Mattis score and x =time from first evaluation). Then, estimated a and b values were used to calculate percent change in Mattis scores over 12 months using the formula: $[(12a/b)*100]$ (Chételat *et al.*, 2005a).

Neuroimaging data acquisition

Within a few days interval at most from inclusion and 18 months later, each patient underwent MRI and ^{18}F FDG-PET scans on the same scanners and using the same acquisition parameters. MRI consisted of a set of 128 adjacent axial cuts parallel to the anterior-posterior commissure (AC-PC) line and with slice thickness 1.5 mm and pixel size 1x1 mm, using the SPGR (spin gradient recalled) sequence (TR=10.3 ms; TE=2.1 ms; FOV=24x18 cm; matrix=256x192). PET data were collected using the ECAT Exact HR+ device with isotropic resolution of 4.2x4.2x4.6 mm (FOV=158 mm). A catheter was introduced in a vein of the arm to inject the radiotracer. Following ^{68}Ga transmission scans, three to five mCi of ^{18}F FDG were injected as a bolus at time 0, and a 10 min PET data acquisition started at 50 min post-injection period. Sixty-three planes were acquired with septa out (volume acquisition), using a voxel size of 2.2x2.2x2.43 mm (see Chételat *et al.*, 2005a for further details).

PET data processing and analysis

To implement the two complementary analyses described in Introduction, metabolic changes were evaluated, first, directly from baseline and follow-up PET data and comparing changes in converters and non-converters. Second, maps reflecting metabolic percent annual changes, called “PET-PAC” maps in what follows, were generated for each patients and used to assess relationships with cognitive decline as well as for supplementary analyses. The following sections will successively briefly describe the common and specific processing steps for these analyses. Further details and illustration of these processing steps are provided as **Supplementary material**.

Common processing steps

A first coregistration was performed to place baseline and follow-up MRI and PET data of each patient in the same space. Second, all PET data were voxel-wise corrected for partial volume effects (PVE) using the patient contemporary MRI and the “modified Muller-Gardner” method (Quarantelli *et al.*, 2004). Third, PET data were scaled using a metabolically preserved brain region, namely the cerebellar vermis (Mevel *et al.*, 2007), to control for inter- and intra-individual global variations in PET signal.

Optimal spatial normalization parameters, to be used in the subsequent specific procedures, were estimated from the spatial normalization of MRI data onto a customized aMCI template using optimal Voxel-Based Morphometry (Good *et al.*, 2001) as previously used in our laboratory (Chételat *et al.*, 2005b). Note that a single set of normalization parameters was estimated for each patient so as to normalize baseline and follow-up PET data using the same parameters to avoid bias due to differential spatial normalization.

PET data processing for analyses comparing baseline and follow-up PET

The optimal normalization parameters were applied to baseline and follow-up coregistered, PVE-corrected and scaled PET data resulting from the common processing steps. Spatially normalized PET data were subsequently smoothed using a Gaussian kernel of 10mm, and entered into the following statistical analyses.

First, a 'Population main effect: 2 cond's, 1 scan/cond' (paired t-test) with 2 conditions (baseline and follow-up) was performed to assess the pattern of metabolic evolution in all aMCI patients by comparing baseline to follow-up data (with a 1 -1 contrast). The resulting SPM-T map was projected onto the aMCI whole brain template.

Second, a 'Multi-group: conditions & covariates' (repeated measures ANOVA) with 2 groups (converters and non-converters) and 2 conditions (baseline and follow-up) was performed on the same data to assess the patterns of metabolic evolution in converters and in non-converters separately, by comparing baseline to follow-up data in each group (with a 1 -1 and a 0 0 1 -1 contrast, respectively). Resulting SPM-T maps were projected onto the customized aMCI whole brain template. To highlight areas of greater metabolic decrease in converters as compared to non-converters, between-group comparison of baseline minus follow-up PET data was then performed (through a 1 -1 -1 1 interaction contrast) onto the voxels exhibiting significant metabolic decreases in converters (using the inclusive masking procedure of SPM2).

Clusters showing significant interaction in the above analysis were also used to define Volumes Of Interest (VOIs) for subsequent application onto PET-PAC maps (see below).

PET data processing for analyses with PET-PAC maps

The baseline and follow-up coregistered, PVE-corrected and scaled PET data resulting from the common processing steps were used to create individual PET-PAC maps. These PET-PAC maps represent the voxel-wise calculation of percent metabolic change over the 18-month follow-up period (i.e. the difference between follow-up and baseline scaled PET value divided by baseline PET value X 100) expressed in annual percent change (i.e. multiplied by 12/18). Note that this calculation was performed only onto those voxels common to both baseline and follow-up PET data, identified using a masking procedure. The optimal spatial normalization parameters were then applied to these PET-PAC maps, which were subsequently smoothed using a Gaussian kernel of 10mm.

A 'Single subjects: covariates only' analysis was then conducted onto these 17 PET-PAC maps using Mattis-PAC as covariate to assess the relationship between metabolism changes and global cognitive decline. Both positive and negative correlations were assessed.

To perform VOI-based discriminant and correlation supplementary analyses, mean PET-PAC values were extracted from each PET-PAC map in the VOIs defined above, using the 'binary ROIs analysis' option of the 'fMRI-ROI analysis' SPM2 toolbox.

To assess the accuracy of the metabolic changes in the VOIs for monitoring the progression to AD in converters, a discriminant analysis was performed. Univariate analyses (T-test) of the mean PET-PAC value of each VOI independently were computed, and a multivariate F-statistic based on MANOVA analysis was performed on all VOIs values combined thanks to a Linear Discriminant.

Finally, to highlight the brain networks whose dysfunction or relative preservation may be related to that of each VOI, correlation analyses were then

performed by entering the mean PET-PAC value of each VOI as covariate in a 'Single subjects: covariates only' voxel-based analysis with PET-PAC maps, assessing both positive and negative correlations, respectively.

All data processing and voxel-based statistical analyses were performed using SPM2 running on MATLAB 6.5. The threshold for significance was set to $p(\text{uncorrected}) < 0.005$, which is identical (Alexander *et al.*, 2002) or more severe (Drzezga *et al.*, 2003) than previously used in longitudinal PET studies in AD and judged to provide the best compromise, neither too permissive nor over-conservative with risk of type 2 errors.

RESULTS

Clinical data

Baseline and follow-up clinical characteristics are presented in **Table 1**. Over the 18 months follow-up period, 7 of the 17 aMCI patients converted to clinical diagnosis of probable AD. Baseline MMSE scores and follow-up Mattis scores were lower in converters than in non-converters. As already reported (Chételat *et al.*, 2005a), the Mattis scores significantly decreased over the follow-up period in converters, but not in non-converters.

Comparison between baseline and follow-up PET

The patterns of metabolic changes from baseline to follow-up in the whole aMCI sample, and in converters and non-converters separately, are illustrated in **Figure 1**. In the whole aMCI sample, metabolic decreases were largely bilateral and involved medially the PCC and frontal areas (Brodmann areas BA 11 and 24/32), and laterally the temporo-parietal cortex (with right predominance), insula and inferior temporal cortex. Assessing converters and non-converters separately, effects were similar but higher in the former and lower in the latter, a difference that was particularly prominent in ventro-medial prefrontal regions (BA25 and 24/32). Also, the PCC changes observed in the whole sample extended to the middle cingulate cortex in converters.

The repeated measures ANOVA comparing converters to non-converters revealed areas of significantly greater metabolism decrease in converters than non-converters, but not in the reverse contrast. These changes were located in two distinct ventro-medial prefrontal regions: the left anterior cingulate cortex (BA24/32)

and the subgenual area (BA25; **Figure 2A**). As described above, these two clusters were then made into VOIs for the correlation and discriminant analyses.

PET-PAC maps

Positive correlation between PET-PAC maps and Mattis-PAC revealed a single significant cluster located in the subgenual area (BA25; **Figure 2B**). The reverse contrast (i.e. PET-PAC increases with Mattis-PAC decreases) disclosed a single cluster located in the right dorso-medial prefrontal cortex (BA9/10; **Figure 3A**).

While a partial overlap was observed between individual values of converters and non-converters using the mean PET-PAC in BA24/32 ($p=0.001$; $AUC=0.94$) or in BA25 ($p=0.006$; $AUC=0.87$) separately, the combination of the mean PET-PAC in these two VOIs improved the between-group discrimination ($p=0.0001$; $AUC=1$; **Figure 4**).

Positive correlation between BA24/32 mean PET-PAC value and PET-PAC maps highlighted surrounding medial prefrontal areas (BA24/32/11) as well as the right PCC including the retrosplenial cortex (BA23/26/29; **Figure 5**). The reverse contrast did not reveal any significant negative correlation.

Positive correlation between BA25 mean PET-PAC value and PET-PAC maps revealed two clusters, the first encompassing surrounding prefrontal areas (BA25/24) and right hippocampus and amygdala, and the second involving the left parahippocampal cortex (BA20; **Figure 5**). The reverse contrast (negative correlation) highlighted two close clusters in the right dorso-medial prefrontal cortex (BA8 and BA9; **Figure 3B**).

DISCUSSION

In the present study, we used a method specifically designed for the longitudinal assessment of PET changes, including voxel-based PVE correction and optimal normalization of each pair of PET data with the same parameters, as well as PET-PAC maps calculation restricted to common GM voxels. This method prevents as far as possible any confounding effects of brain tissue atrophy or methodological bias due to differential normalization and segmentation of baseline and follow-up data. The effects highlighted here are thus thought to reflect genuine metabolic changes taking place during the transition from aMCI to AD.

In the whole aMCI sample, we found progressive metabolic decreases over an 18-month follow-up period encompassing the temporo-parietal cortex and posterior medial parietal areas, consistent with numerous previous studies underlining the early involvement of these areas in AD (see Introduction). Our results also disclosed significant changes in specific prefrontal areas, suggesting that prefrontal metabolic alteration are in fact initiated early in the course of AD. Most notably, the metabolic declines found to be significantly greater in converters relative to non-converters specifically and uniquely pointed to two medial prefrontal areas, namely the anterior cingulate cortex (BA24/32) and the subgenual area (BA25). A similar analysis also pointed to prefrontal areas in Drzezga *et al.* study (2003), but involved lateral prefrontal rather than medial regions. In that study, the medial prefrontal areas were found to show similar decreases in converters and nonconverters which was interpreted as reflecting a normal aging process. Our findings disagree with this interpretation as the two groups did not differ in age or follow-up duration, and furthermore the metabolic changes in both medial prefrontal areas were found not to correlate with age (data not shown). In contradiction with Drzezga *et al.* (2003),

therefore, the present study argues in favour of AD-related pathological processes in these two regions. In support of this contention, the same two medial prefrontal regions have been previously reported to show specific perfusion decreases from the entorhinal to the limbic neuropathologic Braak stages (Braak and Braak, 1991), corresponding respectively to aMCI and early AD (Bradley *et al.*, 2002).

For reasons detailed in Introduction, we also assessed metabolic changes in relation to global cognitive decline across the whole aMCI sample. Positive correlation between PET-PAC maps and Mattis-PAC highlighted a single ventro-medial prefrontal area encompassing the same BA25 region as that found in the between-group comparison, but surprisingly failed to highlight the BA24/32 cluster. As previously proposed (Chételat *et al.*, 2005a), patients expected to present with probable AD criteria at the end of the follow-up period (converters) include both patients with rapid cognitive decline, and patients with less rapid cognitive decline but who started from lower baseline cognitive status. Our findings thus suggest that the metabolic decrease in BA25 is specifically related to the slope of cognitive decline, while that in BA24/32 may instead be related to baseline cognitive performance. Consistent with this hypothesis, we found a significant positive correlation between baseline MMSE performances and BA24/32 PET-PAC values ($p=0.0006$; data not shown), while no significant correlation was found with BA25 PET-PAC values ($p=0.209$; data not shown). Overall, these two regions thus appear to serve complementary roles in expressing the metabolic decreases from aMCI to AD. This was also supported by our multivariate analysis showing improved discrimination between converters and non-converters when combining both BA25 and BA24/32 as compared to either VOI separately. While the complete discrimination found here

would need to be validated from an independent and larger sample, our results strongly support the use of ^{18}F FDG-PET to monitor early AD progression.

So as to better understand the mechanisms underlying these metabolic changes, we also performed metabolic-metabolic correlations between PET-PAC in each VOI and whole brain PET-PAC maps, allowing unravelling the whole brain networks whose metabolic changes relate to those in each of the two prefrontal VOIs (i.e. BA24/32 and BA25). Interestingly, these analyses highlighted two distinct networks for BA24/32 and BA25, the former involving the PCC and the latter the hippocampal region and temporal pole. These distinct relationships suggest that the medial prefrontal metabolic decreases characterizing the progression from aMCI to clinically probable AD may result from disconnection from limbic structures, i.e. from the PCC for BA24/32 and from the hippocampus for BA25. This so-called diaschisis hypothesis (Minoshima *et al.*, 1997; Meguro *et al.*, 2001; Bradley *et al.*, 2002; Chételat *et al.*, 2003b; Nestor *et al.*, 2004) is consistent with recent functional MRI studies of functional connectivity showing, through a method similar to the correlation approach used here, altered hippocampal functional connectivity with the PCC and ventro-medial prefrontal cortex in early AD (Greicius *et al.*, 2004; Wang *et al.*, 2007; Allen *et al.*, 2007). As the uncinate fasciculus directly connects the hippocampus, amygdala and temporal poles to the subgenual cortex (Kier *et al.*, 2004; Schmahmann *et al.*, 2007), disruption of this WM tract may lead to the specific relationships found here. Furthermore, alteration of this tract has been reported in AD (Taoka *et al.*, 2006; Yasmin *et al.*, 2008), and direct hippocampal projection fibers to BA25 were shown to mainly originate from the CA1 subfield (Zhong *et al.*, 2006), i.e. the hippocampal subregion most involved by atrophic processes from aMCI to clinically probable AD (Chételat *et al.*, 2008). The progressive metabolic decrease in

BA25 is thus thought to directly reflect its disconnection from the hippocampus. In contrast, disruption of the rostral cingulum bundle relating the PCC to the frontal cortex (Mufson and Pandya, 1984; Schmahmann *et al.*, 2007; Mori *et al.*, 2008) is probably responsible for the metabolic decrease observed in BA24/32. The caudal part of this tract, which connects the hippocampus to the PCC, is also altered early in AD (Rose *et al.*, 2000; Xie *et al.*, 2005; Medina *et al.*, 2006; Villain *et al.*, 2008) probably accounting for early PCC hypometabolism (Rose *et al.*, 2000; Chételat *et al.*, 2003b; Nestor *et al.*, 2004; Xie *et al.*, 2005; Villain *et al.*, 2008). Our findings suggest that, as aMCI progress to AD, PCC alterations progressively lead to medial prefrontal disruption through involvement of the rostral part of the cingulum bundle. Overall, therefore, BA24/32 metabolic decreases may reflect indirect hippocampo-frontal disconnection processes, as already mentioned elsewhere (Grady *et al.*, 2001; Bradley *et al.*, 2002; Villain *et al.*, 2008) probably mediated by the cingulum bundle which is the major path for fronto-hippocampal connectivity (Kobayashi and Amaral, 2003).

Intriguingly, most structures highlighted in the present study, namely the hippocampus, amygdala, PCC and medial prefrontal cortex, are key components of the episodic memory network (Cabeza and Nyberg, 2000). The role of the uncinate fasciculus and cingulum bundle in memory processes has also been highlighted (Levine *et al.*, 1998; Gaffan and Wilson, 2008; Sepulcre *et al.*, 2008), more specifically for autobiographical memory related to emotional events (Markowitsch *et al.*, 2003). In addition, dysfunction in ventro-medial prefrontal areas has been related to depressive symptoms in healthy subjects (Steele *et al.*, 2007) and apathy in AD (Marshall *et al.*, 2007). Taken together, disruption of the brain networks leading to progressive decrease in ventro-medial prefrontal metabolism may underlie the

worsening of memory impairments as well as the emergence of mood disorders reported as aMCI progresses to clinical AD (Assal and Cummings, 2002).

Finally negative correlations between PET-PAC maps and Mattis-PAC as well as BA25 PET-PAC, both highlighted a single and identical dorso-medial prefrontal region encroaching BA8/9/10. This suggests that, as the disease progresses and BA25 metabolism decreases, BA8/9/10 metabolism relatively increases, potentially reflecting functional compensatory mechanism, as proposed in previous studies for the same dorso-medial prefrontal areas (Grady *et al.*, 2001; Grady *et al.*, 2003; Remy *et al.*, 2005; Wang *et al.*, 2007). The striking difference between metabolic changes taking place in ventro-medial and dorso-medial prefrontal regions, both known to be connected to the hippocampus (Schmahmann *et al.*, 2007) but showing either relative metabolic decreases or increases respectively, would merit further investigations.

In sum, our findings highlight the specific metabolic changes associated with progression from aMCI to clinical AD, showing metabolic decrease in ventro-medial prefrontal BA24/32 and BA25 paralleled by relative increases in dorso-medial BA8/9/10. Prefrontal metabolic disruptions are likely to reflect disconnection from the hippocampus, both indirectly through the posterior cingulate cortex via cingulum bundle breakdown for BA24/32, and directly through uncinate fasciculus disruption for BA25. Metabolic decreases in these two areas combined specifically characterized rapid progression to AD, suggesting the potential of ¹⁸FDG-PET to monitor early AD progression and to test the effects of new therapies.

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Table 1: Demographic and clinical data of aMCI patients at baseline (t0) and at follow-up (t18). ** significant difference between converters and non-converters $p < 0.01$, α significant difference between t0 and t18 $p < 0.05$, $\alpha\alpha$ $p < 0.01$, $\alpha\alpha\alpha$ $p < 0.001$.

Figure 1: Patterns of brain metabolic changes over 18 months in the whole aMCI sample (A), as well as in non-converters (B) and in converters (C) separately, as illustrated by projection of the SPM-T maps onto a 3D representation of the aMCI customized whole brain template.

Figure 2: Brain areas showing significantly greater metabolic decreases in converters compared to non-converters (A-left), used as VOIs represented onto 3D views of the aMCI whole brain template in further analyses (A-right), and significant positive correlation between metabolic decrease (PET-PAC maps) and global cognitive decline (Mattis-PAC – B) as illustrated by SPM-2 ‘Glass brain’ representation and projection of the SPM-T maps (thresholded at $p < 0.005$; $k > 50$ voxels) onto sagittal sections of the aMCI whole brain template. Peak MNI coordinates (xyz), size in voxels (k), and T and P values are indicated for each significant cluster, and correlation plot and R^2 values are also provided.

Figure 3: Brain areas showing significant negative correlation between PET-PAC maps and Mattis-PAC (A) or BA25 PET-PAC value (B) as illustrated by SPM-2 ‘Glass brain’ representations and projection of the SPM-T maps (thresholded at $p < 0.005$; $k > 100$ voxels) onto sagittal section of the aMCI whole brain template. Peak MNI coordinates (xyz), size in voxels (k), and T and P values are indicated for each significant cluster and the corresponding plots and R^2 values are also provided.

Figure 4: Illustration of the discriminant accuracy of the mean PET-PAC values in BA24/32 and BA25 separately, and of both values combined (2D representation) to separate converters from nonconverters.

Figure 5: VOI-based correlation analysis. Brain areas showing significant positive correlation between PET-PAC maps and PET-PAC values in BA25 (**left**), and BA24/32 (**right**), as illustrated in SPM-2 'Glass brain' representations and projection of the SPM-T maps (thresholded at $p < 0.005$; $k > 100$ voxels) onto sagittal sections of the aMCI whole brain template. Peak and sub-peak MNI coordinates (xyz), size in voxels (k), and T and P values are indicated for each significant cluster and the corresponding plots and R^2 values are also provided. **Hcp**: hippocampus.