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# Correlations between plasma and PET beta-amyloid levels in individuals with subjective cognitive decline: the Fundació ACE Healthy Brain Initiative (FACEHBI)

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## Abstract

**Background:** Peripheral biomarkers that identify individuals at risk of developing Alzheimer's disease (AD) or predicting high amyloid beta (A $\beta$ ) brain burden would be highly valuable. To facilitate clinical trials of disease-modifying therapies, plasma concentrations of A $\beta$  species are good candidates for peripheral AD biomarkers, but studies to date have generated conflicting results.

**Methods:** The Fundació ACE Healthy Brain Initiative (FACEHBI) study uses a convenience sample of 200 individuals diagnosed with subjective cognitive decline (SCD) at the Fundació ACE (Barcelona, Spain) who underwent amyloid florbetaben (<sup>18</sup>F) (FBB) positron emission tomography (PET) brain imaging. Baseline plasma samples from FACEHBI subjects (aged 65.9  $\pm$  7.2 years) were analyzed using the ABtest (Araclon Biotech). This test directly determines the free plasma (FP) and total plasma (TP) levels of A $\beta$ 40 and A $\beta$ 42 peptides. The association between A $\beta$ 40 and A $\beta$ 42 plasma levels and FBB-PET global standardized uptake value ratio (SUVR) was determined using correlations and linear regression-based methods. The effect of the *APOE* genotype on plasma A $\beta$  levels and FBB-PET was also assessed. Finally, various models including different combinations of demographics, genetics, and A $\beta$  plasma levels were constructed using logistic regression and area under the receiver operating characteristic curve (AUROC) analyses to evaluate their ability for discriminating which subjects presented brain amyloidosis.

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**Results:** FBB-PET global SUVR correlated weakly but significantly with A $\beta$ 42/40 plasma ratios. For TP42/40, this observation persisted after controlling for age and *APOE*  $\epsilon$ 4 allele carrier status ( $R^2 = 0.193$ ,  $p = 1.01E-09$ ). The ROC curve demonstrated that plasma A $\beta$  measurements are not superior to *APOE* and age in combination in predicting brain amyloidosis. It is noteworthy that using a simple preselection tool (the TP42/40 ratio with an empirical cut-off value of 0.08) optimizes the sensitivity and reduces the number of individuals subjected to A $\beta$  FBB-PET scanners to 52.8%. No significant dependency was observed between *APOE* genotype and plasma A $\beta$  measurements ( $p$  value for interaction = 0.105).

**Conclusion:** Brain and plasma A $\beta$  levels are partially correlated in individuals diagnosed with SCD. A $\beta$  plasma measurements, particularly the TP42/40 ratio, could generate a new recruitment strategy independent of the *APOE* genotype that would improve identification of SCD subjects with brain amyloidosis and reduce the rate of screening failures in preclinical AD studies. Independent replication of these findings is warranted.

**Keywords:** Subjective cognitive decline, Preclinical AD, Alzheimer's disease, Amyloid  $\beta$ , Plasma biomarker, TP42/40, PET, Florbetaben

## Highlights

- Brain and plasma A $\beta$  levels are partially correlated in SCD subjects.
- Plasma A $\beta$  measurements are independent of *APOE* genotype.
- The model including only plasma TP42/40 level as a variable achieved the highest sensitivity in predicting A $\beta$  PET positivity (83%).
- A simple preselection step using the TP42/40 classifier with an empirical cut-off value of 0.08 would reduce the number of individuals subjected to A $\beta$  FBB-PET by 52.8%.

## Background

Alzheimer's disease (AD), the most common cause of dementia, is a neurodegenerative disorder characterized by progressive memory loss and cognitive decline [1]. Pathological findings of AD include deposits of amyloid beta (A $\beta$ ) peptides in the brain conforming extracellular amyloid plaques together with intracellular deposits of hyperphosphorylated tau [2]. The progressive increase of both pathological hallmarks is associated with gradual synaptic and neuronal loss resulting in the clinical deterioration of patients [3].

There are no effective disease-modifying therapies for AD available at the current time. Neuropsychological assessment [4], cerebrospinal fluid [5] (CSF) analysis, and amyloid positron emission tomography (PET) scans are common methods used for prodromal AD detection. CSF and amyloid PET provide the most reliable in-vivo biomarkers of prodromal AD, but they are not suitable for population screening purposes due to the invasive CSF sampling procedure and the high cost and limited availability of amyloid PET imaging [6, 7]. Magnetic resonance imaging (MRI)-based AD biomarkers have demonstrated high sensitivity to prodromal AD [8]; however, the specificity of MRI is limited for predicting conversion of

mild cognitive impairment (MCI) to dementia [9] and MRI is also impractical in patients with some types of pacemakers, metal implants, or claustrophobia. Consequently, despite the robustness of these biomarkers, they are not suitable for broad population screening in primary care clinical settings. Therefore, there is a growing need for accurate identification of asymptomatic (preclinical) individuals with underlying AD pathology to improve diagnosis and subject inclusion in prevention trials of prodromal and presymptomatic AD.

Discovery of blood-based AD biomarkers would entail important cost-benefit and scalability advantages over current techniques, potentially enabling broader clinical access and efficient population screening. The plasma concentration of A $\beta$  is a logical candidate, but studies to date have produced conflicting results on its utility [10]. Several longitudinal studies with large cohorts such as the Framingham Study [11] with 2189 dementia-free participants followed from baseline until they developed dementia, died, or had been followed for 10 years and the Rotterdam Study [12] with 1756 participants and 392 incident dementia cases identified (follow-up mean 8.6 years) have reported increased risk of dementia associated with lower A $\beta$ 42/40 plasma ratios and that a reduction in plasma A $\beta$ 42 levels over time is linked with cognitive decline [13, 14]. A recent publication [15] studied the ability of A $\beta$  precursor protein (APP/A $\beta$ 42), A $\beta$ 40/A $\beta$ 42 ratios, and their composites to predict individual brain A $\beta^{+/-}$  status determined by A $\beta$ -PET imaging. The results showed that all test biomarkers correlated with both A $\beta$  PET burden and levels of A $\beta$ 42 in CSF in two independent cohorts, demonstrating that the three different types of A $\beta$ -related biomarkers (plasma, CSF, and PET imaging) are highly correlated with each other, clearly indicating the potential utility of plasma biomarkers. Furthermore, an independent study [16] suggests that individuals with subjective cognitive decline (SCD) exhibit significantly higher A $\beta$ 42 plasma concentrations compared with participants

with no complaints. However, other studies have reported a weak or even a lack of association of plasma A $\beta$ 42/40 ratio with AD diagnosis [17–19].

Given that both subjective complaints and impaired episodic memory are present in MCI, the existence of an earlier distinct clinical stage where subjective complaints exist in the absence of detectable objective cognitive deficits is plausible [20]. There is evidence suggesting that SCD may increase the risk of progression to cognitive impairment and dementia [21], and that individuals with SCD have a higher risk of developing AD [22], and present more functional deficits [23] and AD brain pathology than non-SCD participants [24]. SCD might represent the earliest point on the continuum of clinical Alzheimer's symptomatology [25–27], even anticipating the onset of subtle but detectable neuropsychological or biological alterations. Hence, a better understanding of the baseline characteristics of this group of patients may enhance our knowledge of early AD processes, facilitating early diagnosis, follow-up, and preventive treatment, making SCD an interesting target population to study.

The primary aim of this study was to assess the association between plasma A $\beta$  levels and amyloid brain burden. Specifically, we measured A $\beta$ 42 and A $\beta$ 40 plasma levels using two specific sandwich enzyme-linked immunosorbent assay (ELISA) kits, ABtest40 and ABtest42 (Araclon Biotech, Zaragoza, Spain), and quantified amyloid brain burden using florbetaben (<sup>18</sup>F) (FBB)-PET global standardized uptake value ratio (SUVR) in 200 individuals with SCD. We evaluated whether plasma A $\beta$  ratios may be useful biomarkers for AD and a screening tool for amyloidosis in healthy populations.

## Methods

### The FACEHBI cohort

The Fundació ACE Healthy Brain Initiative (FACEHBI) uses a convenience sample of 200 individuals (mean age 65.8  $\pm$  7.2 years; 37.5% males) diagnosed with SCD at Fundació ACE (Barcelona, Spain) recruited from Open House initiatives [28]. The cohort comprised of 52 (26%) APOE  $\epsilon$ 4 allele carriers and 18 (9%) individuals with a positive (SUVR > 1.45) FBB-PET scan. The demographic characteristics of the study cohort are summarized in Table 1 and Additional file 1 (Table S1) by FBB-PET status.

The SCD criteria used to recruit subjects in this study have been described previously [29]. Briefly, inclusion criteria were: 1) subjective cognitive complaints defined as a score of  $\geq$  8 on MFE-30, the Spanish version of the Memory Failures in Everyday Life Questionnaire [30]; 2) Mini-Mental State Examination (MMSE) score  $\geq$  27; 3) Clinical Dementia Rating (CDR) = 0; and 4) performance on the Fundació ACE Neuropsychological Battery (NBACE) [31] within the normal range for age and educational level.

**Table 1** Demographics and clinical characteristics of the study cohort (FACEHBI [29])

Variable	SCD
Subjects, <i>n</i>	200
Age, years	65.87 (7.23)
Education, years	14.76 (4.73)
Gender, % males	37.5
APOE, % $\epsilon$ 4 allele carriers	26
Creatinine, mg/dl	0.92 (0.15)
Body mass index, kg/m <sup>2</sup>	26.64 (4.32)
Hematocrit, %	43.15 (4.93)
FBB-PET SUVR	1.2 (0.15)
FP42/40	0.04 (0.03)
TP42/40	0.09 (0.06)
FP40/TP40	0.44 (0.06)
BP42/40	0.13 (0.09)
FP42/TP42	0.24 (0.21)

Data are shown as mean (SD) unless otherwise specified

APOE apolipoprotein, BP bound peptide, FBB florbetaben (<sup>18</sup>F), FP free plasma, PET positron emission tomography, SCD subjective cognitive decline, SUVR standardized uptake value ratio, TP total plasma

Exclusion criteria were as follows: 1) relevant symptoms of anxiety or depression defined as a score of  $\geq$  11 on the Hospital Anxiety and Depression Scale (HADS) [32]; 2) presence of other psychiatric diagnosis; 3) history of alcoholism and epilepsy; and 4) known renal or liver failure.

Cognitive assessment was performed according to the routines of the Memory Clinic of Fundació ACE as described elsewhere [33]. Baseline MRI of these subjects demonstrated the absence of signs indicative of brain pathology. All participants gave written consent and the protocol was approved by the ethics committee of the Hospital Clinic i Provincial (Barcelona, Spain) (EudraCT: 2014–000798-38).

### MRI acquisition

All MRI scans were acquired prior to FBB-PET. MRI were performed on a 1.5-T Siemens Magnetom Aera (Erlangen, Germany) using a 32-channel head coil. Anatomical T1-weighted images were acquired using a rapid acquisition gradient-echo three-dimensional (3D) magnetization-prepared rapid gradient-echo (MPRAGE) sequence with the following parameters: repetition time (TR) 2.200 ms, echo time (TE) 2.66 ms, inversion time (TI) 900 ms, flip angle 8°, field of view (FOV) 250 mm, slice thickness 1 mm, and isotropic voxel size 1  $\times$  1  $\times$  1 mm. Subjects also received axial T2-weighted, 3D isotropic fast fluid-attenuated inversion recovery (FLAIR) and axial T2\*-weighted sequences to detect significant vascular pathology or microbleeds.

### FBB-PET acquisition

FBB-PET scans were obtained with a Siemens© Biograph molecular-CT machine. PET images were acquired in 20 min starting from 90 min after intravenous administration of 300 Mbq of Florbetaben ( $^{18}\text{F}$ ) radio tracer (NeuraCeq©), administered as a single slow intravenous bolus (6 s/ml) in a total volume of up to 10 ml.

### SUVR estimation

MRI cortical [34] and subcortical [35] parcellations were carried out with Freesurfer 5.3 (<http://surfer.nmr.mgh.harvard.edu/>), following the pipeline described in <https://surfer.nmr.mgh.harvard.edu/fswiki/recon-all>.

FBB-PET were coregistered to the MRI labeled data with the FSL 5.0 software package (<https://fsl.fmrib.ox.ac.uk/fsl/fslwiki>) by means of MCFLIRT, it is an intra-modal motion correction tool based on optimization and registration techniques from FLIRT (FMRIB's Linear Image Registration Tool), which next was also used. These are fully automated tools implemented in FSL 5.0 for linear (affine) intra- and inter-modal brain image registration [36, 37].

Amyloid cortical SUVR was determined as the average of the standardized uptake value normalized by the uptake in the cerebellar grey matter, with this reference region being selected from the MRI cerebellum external and cortex segments. Based on previous studies [38], a cut-off for SUVR above or equal to 1.45 was selected as the amyloid positivity criterion.

### Blood sampling, APOE genotyping, biochemical determinations, and A $\beta$ measurements

Blood samples and the *APOE* genotype from each participant were routinely processed in Fundació ACE as previously described [29, 39]. In brief, blood samples were obtained in the morning after an overnight fast, collected in polypropylene vials with ethylenediaminetetraacetic acid (EDTA) and immediately refrigerated. Samples were centrifuged within 24 h from extraction to collect the plasma and then aliquoted and frozen at  $-80\text{ }^{\circ}\text{C}$  until assayed. Biochemical and hematologic measurements were determined in a reference laboratory according to clinical standards.

For plasma amyloid testing, four determinations were made (Additional file 2). Total plasma (TP) and free plasma (FP) A $\beta$ 40 and A $\beta$ 42 levels were quantified using two specific sandwich ELISA kits, A $\beta$ test40 and A $\beta$ test42 (Araclon Biotech, Zaragoza, Spain), in accordance with the manufacturer's instructions as described elsewhere [39]. Briefly, before analysis, each plasma sample was split into two aliquots: an undiluted aliquot and another aliquot pretreated by 1:3 dilutions in a formulated sample buffer (phosphate-buffered saline (PBS) 0.5 M, 0.5% Tween-20, 1% blocking polymer) intended to break A $\beta$  interactions with other plasma components. Thus, levels of free and total A $\beta$ 40 and A $\beta$ 42 were separately determined

in undiluted and diluted plasma, respectively. The difference between TP and FP concentration corresponds to the amount of amyloid peptide bound to plasma components (BP). The A $\beta$ 42/A $\beta$ 40 ratios in each of these plasma fractions (FP42/40, TP42/40, BP42/40, FP40/TP40, and FP42/TP42) were calculated and served as the target plasma biomarkers for this study.

The levels of TP and FP obtained from plasma samples were expressed as picograms (pg) of A $\beta$  peptide per milliliter (ml) of plasma. The analyses were always performed in duplicates of the same aliquot and in a coded manner to ensure blindness of the operator.

Both inter-assay and intra-assay coefficients of variation were below 5% and 8–20% for ABtest40 and ABtest42, respectively. The detection limit was 3.13 and 200 pg/ml for ABtest40 and 1.56 and 100 pg/ml for ABtest42. One sample was removed from the original FACEHBI cohort [29] because both ABtest determinations were outside the upper limit of quantification ( $> \text{ULQ}$ ). In ABtest42, 84 of 400 (21%) determinations were also outside the quantification range, either because they were below the lower limit of quantification ( $< \text{LLQ}$ ) or due to undetectable peptide levels. We assigned the minimum value of quantification (1.56 pg/ml) to these samples.

### Statistical analysis

We performed several correlation and regression analyses to explore the association between plasma amyloid ratios and FBB-PET brain amyloid burden. First, we conducted a linear regression analysis using FBB-PET global SUVR as the quantitative response variable in SCD subjects. FBB-PET global SUVR was log-transformed for all analyses since it was not normally distributed. The distribution of variables and Shapiro-Wilk test are given in Additional file 3 (Figure S1). We conducted an exploratory analysis with three different transformations for the plasma A $\beta$ 42/A $\beta$ 40 ratios: dichotomous (with regard to the median of the population), quartile, and logarithmic. First, we performed Pearson and Spearman correlation analyses between log-transformed FBB-PET global SUVR and the raw values of each plasma A $\beta$  measure of interest as well as the transformed plasma A $\beta$  ratios (Table 2 and Additional file 4: Table S2). Next, we performed a linear regression analysis using a backward-selection procedure with FBB-PET global SUVR as the quantitative dependent variable, with age, gender, education, *APOE*  $\epsilon$ 4 carrier status, and the best performing log-transformed plasma A $\beta$ 42/40 ratio as independent variables (Table 3 and Additional file 5: Table S3). Bonferroni correction was used to adjust for multiple comparisons.

We used logistic regression to construct four different models (Table 4) to evaluate the usefulness of the covariates selected from the backward regression model for discriminating which SCD participants were FBB-PET

**Table 2** Correlation between direct A $\beta$  plasma and log-transformed FBB-PET SUVR

Logarithmic	FP42/40	TP42/40	FP42/TP42	FP40/TP40
L_PET				
Pearson's $r$ ( $n = 199$ )	-0.160*	-0.248**	0.100	0.085
$p$ value (2-tailed)	0.024	4.04E-04	0.162	0.231
95% confidence interval	-0.292 to -0.021	-0.374 to -0.113	-0.04 to 0.236	-0.055 to 0.221

Plasma amyloid beta (A $\beta$ )42/40 ratios were transformed in logarithmic scale

Bonferroni correction was used to adjust for multiple comparisons ( $< 1.92E-03$ )

FP free plasma, L\_PET, logarithmic transformed positron emission tomography score, TP total plasma

\* $p \leq 0.05$

\*\* $p \leq 0.01$

amyloid positive ( $> 1.45$ ) in 199 participants. The models were structured to reflect categories of predictive information by the ease of its acquisition. Accordingly, the first model (model #1) included only predictors that can be easily obtained (age). The second model additionally requires a blood extraction and includes two parts: model #2a for *APOE*  $\epsilon 4$  carrier status (0–1) which served as the reference model for discrimination of amyloid PET-positive subjects as proposed by Petersen [25], and model #2b for a plasma determination of TP42/40 in log units. The third model (model #3) included the three variables described above (age, *APOE*, and TP42/40). Finally, the fourth model (model #4) only included the target plasma biomarker (logTP42/40). We used the area under the receiver operating characteristic curve (AUROC) from the models as a measure of how well the model discriminated between FBB-PET positive and negative subjects. The criterion for choosing the operating point along the ROC curve was Youden's index maximum. The logistic models allowed us to assign a predicted probability of being FBB-PET SUVR positive to each subject based on values for the selected variables in the model. In addition to sensitivity/specificity performance measures, the predictive values (positive (PPV) and negative (NPV)) of the models were calculated.

Finally, the effect of *APOE* genotype on plasma A $\beta$  levels was assessed by comparing A $\beta$  plasma measurements between *APOE*  $\epsilon 4$  carriers and noncarriers by analysis

**Table 3** Backward selection regression analysis: amyloid beta plasma TP42/40 ratio and log FBB-PET global SUVR with covariates

	Estimate	Standard error	T value	$p$ value
(Intercept)	-0.0701	0.030	-2.301	0.022*
Age	0.0015	4.45E-04	3.573	4.45E-04***
<i>APOE</i>	0.035	0.007	5.107	7.75E-07***
Log TP42/40	-0.041	0.011	-3.794	1.98E-04***

Residual standard error = 0.042 on 195 degrees of freedom (DF)

Adjusted  $R^2 = 0.193$ ;  $F = 16.75$  on 3 and 195 DF;  $p$  value = 1.01E-09

Backward selection regression analysis adjusting for age, *APOE* and TP42/40; statistical significance was set to  $p < 1.92E-03$  after Bonferroni correction for multiple comparisons

*APOE* apolipoprotein, FBB florbetaben ( $^{18}F$ ), PET positron emission tomography, SUVR standardized uptake value ratio, TP total plasma

\* $p \leq 0.05$

\*\*\* $p < 0.001$

of variance (ANOVA) (Additional file 6: Table S4) by performing separate regression analyses between logTP42/40 and FBB-PET global SUVR in *APOE*  $\epsilon 4$  carriers and noncarriers (Additional file 7: Figure S4), and by testing the interaction term between *APOE*  $\epsilon 4$  carrier status and logTP42/40 in the logistic regression model #3 described above. Statistical analysis was performed with SPSS 19 and RStudio Version 1.0.136. The Ggplot2 package was used for graphic representations.

## Results

### Relationship between A $\beta$ plasma ratio and FBB-PET

The FACEHBI study has been designed to identify the most important factors related to preclinical AD [29]. To evaluate the strength of the association between plasma amyloid biomarkers and A $\beta$ -PET burden, we conducted correlation analyses. Logarithmic TP42/40 and FP42/40 showed significant negative Pearson's correlations with amyloid PET burden, although only TP42/40 exceeds the Bonferroni correction ( $r = -0.248$  (-0.374 to -0.113);  $p = 4.04E-04$ ). In contrast, direct plasma levels of A $\beta 40$  and A $\beta 42$  did not significantly correlate with FBB-PET global SUVR (Additional file 4: Table S2C). BP42/40 was excluded from further analyses due to collinearity with TP42/40 (Pearson's  $r = 0.972$  (0.963–0.979);  $p < 2.2E-16$ ; Additional file 4: Table S2A).

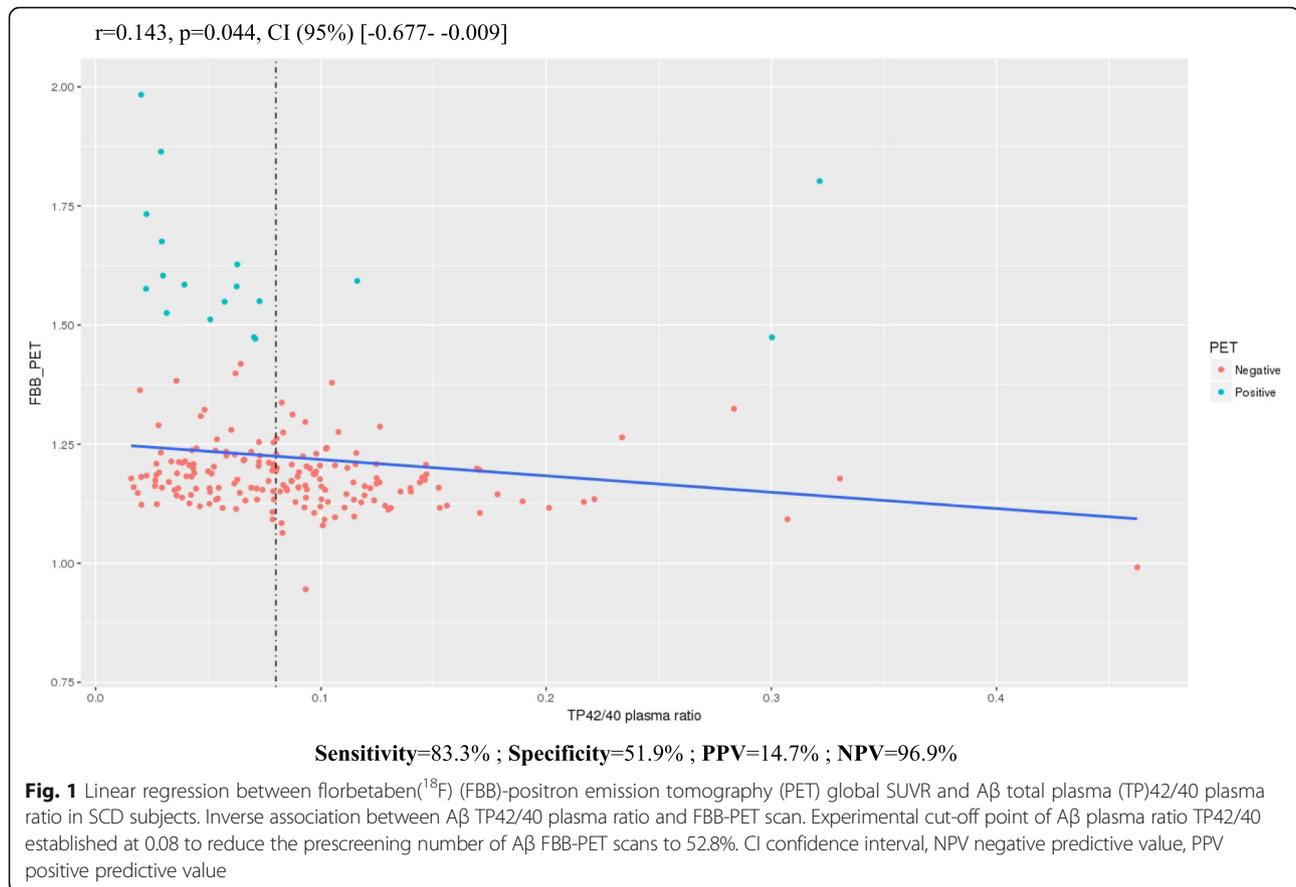
Backward regression analysis identified age, *APOE*  $\epsilon 4$  status (0–1), and logTP42/40 as significant covariates of the best model predicting FBB-PET global SUVR ( $R^2 = 0.193$  and  $p$  value = 1.01E-09; Table 3). The inverse association between FBB-PET SUVR and TP42/40 is graphically represented with raw data in Fig. 1. The associations with the other A $\beta$  plasma biomarkers are shown in Additional file 8 (Figure S2). After stratifying for *APOE*  $\epsilon 4$ , the linear regression analysis showed a negative relationship between plasma TP42/40 and FBB-PET uptake ( $r = -0.523$  (-0.185 to -0.067);  $p = 8.12E-05$ ) exclusively in *APOE*  $\epsilon 4$  carriers (Additional file 9: Figure S3).

To assess the relevance of the plasma biomarkers in predicting amyloid PET positivity, the TP42/40 model was selected for the subsequent AUROC analysis. Education and gender were excluded due to their lack of significance

**Table 4** Summary of logistic regression models and AUROC analysis

Characteristic	Model 1		Model 2a		Model 2b		Model 3		Model 4	
	OR (95% CI)	p	OR (95% CI)	p	OR (95% CI)	p	OR (95% CI)	p	OR (95% CI)	p
Age	1.091 (1.016–1.172)	0.017	1.114 (1.033–1.202)	0.005	1.097 (1.017–1.183)	0.017	1.113 (1.029–1.205)	0.008	–	–
APOE	–	–	7.319 (2.503–21.401)	2.8E-04	–	–	7.208 (2.431–21.373)	3.7E-4	–	–
Log TP42/40	–	–	–	–	0.131 (0.021–0.819)	0.030	0.126 (0.017–0.944)	0.044	0.133 (0.021–0.829)	0.031
AUROC	0.702		0.806		0.754		0.818		0.681	
Youden's index	0.42		0.54		0.49		0.55		0.42	
Cut-off	0.101		0.147		0.115		0.092		0.081	
Specificity	69.6		87.3		76.8		77.3		59.1	
Sensitivity	72.2		66.7		72.2		77.8		83.3	
PPV	19.1		34.3		23.6		25.5		16.7	
NPV	96.2		96.3		96.5		97.2		97.2	

Logistic regression models were used to assess predictors of FBB-PET SUVR positivity (cut-off > 1.45) after adjustment by selected covariates in 199 participants. Model 1 included only age as a predictor; model 2 requires a blood extraction (APOE ε4 carrier status (0–1) or log TP42/40, model 2a and 2b respectively); model 3 included age, APOE, and log TP42/40, and model 4 only included the target plasma biomarker log TP42/40. The criterion for choosing the operating point along the ROC curve was Youden's index maximum. AUROC area under the receiver operating characteristic curve, CI confidence interval, NPV negative predictive value, OR odds ratio, PPV positive predictive value, TP total plasma



in the backward regression model. When A $\beta$ -PET was used as the standard classifier for A $\beta^+$ /A $\beta^-$  status, all models worked in a similar way to the reference discrimination model #2a with age and *APOE* as predictors (AUROCs of 0.702, 0.806, 0.754, 0.818, and 0.681 for models #1, #2a, #2b, #3, and #4 respectively; Table 4, Fig. 2). The effect and significance of TP42/40 was maintained in the different models indicating a robust association with A $\beta$ -PET positivity. Model #2a presented the best balance between PPV/NPV (34.3–96.3%, respectively), but at the same time showed the lowest sensitivity (66.7%). On the other hand, TP42/40 alone (model #4) achieved the best sensitivity (83.3%) and a good NPV (97.2%), indicating its value as a potential screening tool for detecting brain amyloidosis (Table 4). Using an empirical cut-off point of TP42/40 = 0.08, individuals with a TP42/40 plasma ratio < 0.08 (52.8%) would be prescreened with a FBB-PET scan, capturing 83% of the positive amyloid cases, thus reducing the prescreening number of A $\beta$  FBB-PET (sensitivity = 83.3%; specificity = 51.9%; NPV = 96.9%; PPV = 14.7%; Fig. 1).

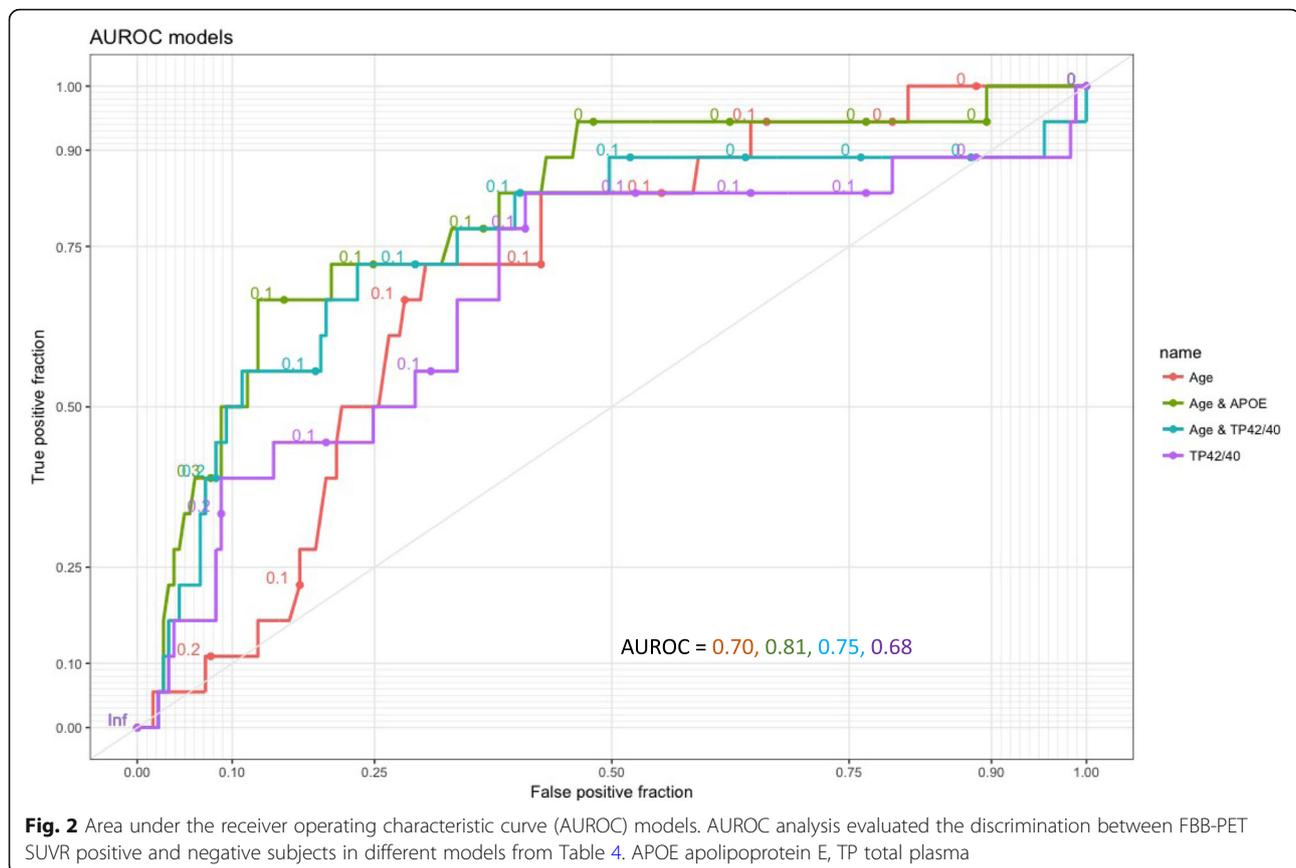
#### Effect of *APOE* genotype on plasma A $\beta$ levels

In the current study, we found no association between *APOE* genotype and plasma A $\beta$  measurements, indicating

independence between both variables. No plasma A $\beta$  measure significantly differed between *APOE*  $\epsilon 4$  carriers and noncarriers (Additional file 6: Table S4 and Additional file 7: Figure 4). This independence, confirmed by the absence of significance for the interaction term between *APOE*  $\epsilon 4$  and logTP42/40 (Additional file 10: Table S5) (odds ratio (OR) 0.022, 95% confidence interval (CI) 2.30E-04 to 2.201);  $p = 0.105$ ) could be an advantage if using this biomarker as a screening tool since it would avoid bias resulting from *APOE* screening.

#### Discussion

The FACEHBI study has been designed to identify the most relevant factors related to preclinical AD in a cohort of individuals with SCD [29]. FACEHBI has a 9% prevalence of amyloid PET positivity, which is lower than similar series reported in the literature. Ossenkoppele et al. [40] estimated a prevalence of 11% brain amyloid positivity in a cohort of healthy controls aged 55–64 years, and 22% in those aged 65–74 years. In a meta-analysis [41], Jansen et al. reported a prevalence of amyloid PET positivity of approximately 20% at age 65 years. The Mayo Clinic population study [42] showed a prevalence of amyloid PET positivity of 13% in the age group 60–64 years and 32% in those aged 65 to 69 years. A possible cause for the low



prevalence of amyloid PET positivity in the FACEHBI cohort is that a strict definition of cognitive normality was used. A score of 1.5 SD below the mean according to age and level of education in any single NBACE [43] test precluded individuals from enrolling into the FACEHBI study. Other studies with a more liberal definition of cognitive normality included patients that would have been considered to have MCI by our standards, presumably increasing their prevalence of amyloid PET positivity. Secondly, the setting of the study is relevant, as it is well known that participants from clinical samples tend to show higher risk of cognitive progression (and probably greater brain amyloidosis) than those from population-based samples and healthy volunteers, even though both groups are considered to be cognitively normal. In this regard, FACEHBI is a mixed sample, but most of our participants (70%) are healthy volunteers from the community who came to check their cognition for free through Open Door Initiatives. This could partly explain a lower prevalence of brain amyloidosis in our FACEHBI participants compared with pure clinical samples.

The main finding of this study is that lower plasma A $\beta$ 42/40 ratios (particularly the TP42/40 ratio) correlate with higher cerebral A $\beta$  plaque burden assessed by amyloid FBB-PET imaging in the FACEHBI SCD cohort. This

inverse correlation is presumably driven by the reduction of A $\beta$ 42 and the increase of A $\beta$ 40 in the A $\beta$ <sup>+</sup> population (Additional file 11: Figure S5). These results are independent of previous explorations and are in line with other promising results reporting similar associations between plasma A $\beta$ 42/40 ratio and cortical fibrillary A $\beta$  burden [15, 44–50] (for review see [51–53]). This study provides added value as it is one of few [48, 49] that explores the association between A $\beta$  plasma ratios and A $\beta$  brain burden within a population of cognitively normal individuals, avoiding the possibility of potential circular associations related to inclusion of MCI and AD subjects along with healthy controls in the same models. Nevertheless, discrepant results from other studies [17–19, 54–56] that assessed the performance of plasma A $\beta$  levels in predicting the A $\beta$  brain status cannot be disregarded. Part of this controversy could be explained by the mixed distribution of individuals with and without cerebral A $\beta$  deposition (as quantified by amyloid PET and/or by CSF analysis) among healthy controls, MCI, and demented individuals.

It is believed that the clearance of brain A $\beta$  is reduced in AD patients compared with healthy controls. This is consistent with a report by Giedraitis et al. [57] who found a correlation between CSF and plasma levels of both A $\beta$ 40 and A $\beta$ 42 in healthy individuals, whereas no

correlations were seen in AD or MCI patients. Thus, the search for an association between blood and brain A $\beta$  levels should be directed towards the earliest stages of the disease (preclinical/prodromal AD), which is also when it is of maximum clinical interest especially as a target population for the development of novel disease-modifying therapies. However, it has been difficult to draw definite conclusions with respect to changes in plasma A $\beta$  concentration in AD [52] because of the inconsistency of the available data. Stringent standardization is required to obtain reliable data that facilitate comparison between studies. In this study we used A $\beta$ 42/A $\beta$ 40 plasma ratios (particularly the TP42/40 ratio) instead of single peptide measurements to attenuate possible bias in single A $\beta$  peptide level quantifications caused by inconsistencies in sample handling [58].

The regression model that included only the TP42/40 ratio did not show sufficient predictive ability to identify those individuals with a positive FBB-PET scan, accounting for only 20% of the variance. Clearly, screening with these factors would not be an acceptable option for determining amyloid PET positivity in the clinical practice setting. Nevertheless, the plasma TP42/40 ratio showed a significant negative correlation with FBB-PET SUVR. This suggests that this plasma A $\beta$  biomarker could be useful as an enrichment tool to identify potential candidates for clinical trials focused on preclinical AD. To prove this, we would need to reproduce the results in a controlled trial with an independent sample. Our analyses suggest that inclusion of the TP42/40 plasma biomarker in a classifier model could reduce unnecessary amyloid PET scans, facilitating recruitment for clinical trials. Taking this into account, in a clinical trial recruiting scenario targeting cognitively normal people, a prescreening step using a TP42/40 classifier (cut-off value = 0.08) would reduce the number of individuals undergoing A $\beta$  FBB-PET scans to 52.8%. The cortical A $\beta$  burden of these subjects would have to then be confirmed by A $\beta$  FBB-PET scans. Consequently, this strategy would reduce the costs [59] of identifying individuals with brain amyloidosis for AD prevention trials [60].

We observed an association of age with plasma A $\beta$  ratios as described in previous studies [41, 42, 59, 61]. No association was found between the *APOE*  $\epsilon$ 4 genotype and A $\beta$  plasma ratios, demonstrating independence between *APOE*  $\epsilon$ 4 genotype and this candidate plasma biomarker. The linear regression analysis stratifying for *APOE*  $\epsilon$ 4 showed a negative relationship between TP42/40 and FBB-PET SUVR in *APOE*  $\epsilon$ 4 carriers but not in noncarrier SCD individuals. At first glance, these results seem contradictory with other studies reporting a significant negative relationship between plasma A $\beta$  and amyloid PET only in *APOE*  $\epsilon$ 4 noncarriers [46, 48, 62, 63]. One possible explanation could stem from the difference in cohort composition, as the previous studies included patients

with MCI and AD diagnosis, while our sample is comprised only of SCD individuals. Therefore, their *APOE*  $\epsilon$ 4 carrier group included participants who were older and more cognitively impaired than ours, whereas their *APOE*  $\epsilon$ 4 noncarrier group could be more similar to our *APOE*  $\epsilon$ 4 carrier group in terms of demographics and cognition. Therefore, they observed a negative correlation between A $\beta$  plasma and PET in *APOE*  $\epsilon$ 4 noncarriers that would be equivalent to the correlation in *APOE*  $\epsilon$ 4 carriers in our study. We attribute this finding to the potential enrichment of preclinical AD cases in the *APOE*  $\epsilon$ 4<sup>+</sup> SCD subgroup. Specifically, our hypothesis is that the rate of genuine AD cases contained in a study population might distort the correlation between A $\beta$ -PET and plasma amyloid measurements.

We consider one of the main strengths of this study is that it includes a well-defined homogeneous population putatively positioned at a very early stage of the disease. We know that the main risk factors such as age and *APOE* do not follow the correlation expected in advanced stages of AD [64], and we have previously reported [64] that the *APOE*  $\epsilon$ 4 genotype had significant effects on the association with FBB-PET global SUVR in SCD subjects. Thus, AD does not behave linearly, and it could be that the TP42/40 ratio behaves independently from *APOE* when positioned to the left of the disease continuum. Our data show that refraining from strict inclusion criteria, such as *APOE*  $\epsilon$ 4 positivity, will be important to avoid detection bias.

An important limitation of this study is the fact that the FBB-PET cut-off value for positivity is arbitrary in SCD populations. The global SUVR > 1.45 cut-off value has been calculated for dementia patients but perhaps it should be adjusted for populations with different degrees of cognitive impairment or even on different segments of the AD continuum. Another limitation is the small sample size which warrants independent replication. Although Fandos et al. [49] reported similar results from the AIBL dataset in cognitively healthy and SCD individuals [65], it would be interesting to repeat the same analysis by A $\beta$  cluster and replicate our findings in a larger population with a higher rate of amyloid PET-positive individuals to improve discrimination and accuracy of the plasma amyloid cut-off point.

Future research should address whether the association between brain and plasma A $\beta$  levels in SCD participants is able to discriminate those older adults who will experience a fast cognitive decline from those who will remain stable over time.

## Conclusion

The present data show an inverse association between plasma A $\beta$ 42/40 ratios and brain fibrillary A $\beta$  deposition in SCD participants. Including the TP42/40 plasma ratio

could help generate a more cost-effective recruitment strategy for clinical trials independent of the *APOE* genotype (reflecting the real diversity of the *APOE* genotype in preclinical AD) and reducing the associated costs of preselecting subjects using expensive imaging techniques.

## Additional files

**Additional file 1: Table S1.** Demographics and clinical characteristics of subjects studied (FACEHBI [29]) for FBB-PET status being positive > 1.45. (DOCX 31 kb)

**Additional file 2:** Plasma amyloid beta levels measured with ABtest for the FACEHBI samples. (XLSX 28 kb)

**Additional file 3: Figure S1.** A) Distribution of FBB-PET and plasma ratios. B) Shapiro-Wilk test for FBB-PET and plasma ratios. C) Log distributions FBB-PET and plasma ratios. D) Shapiro-Wilk test for logarithmic FBB-PET and log-plasma ratios. A, B) Distributions and Shapiro-Wilk test for plasma ratios and FBB-PET to test normality. C, D) Distributions and Shapiro-Wilk test for transformed to logarithmic plasma ratios and FBB-PET to test normality. (PDF 299 kb)

**Additional file 4: Table S2.** Exploratory analysis. (DOCX 23 kb)

**Additional file 5: Table S3.** Regression analyses between A $\beta$  plasma ratios and FBB-PET SUVR. (DOCX 18 kb)

**Additional file 6: Table S4.** ANOVAs comparing *APOE*  $\epsilon$ 4 carriers vs noncarriers. (DOCX 14 kb)

**Additional file 7: Figure S4.** *APOE* and plasma A $\beta$  ratios. The effects of *APOE* genotype on plasma A $\beta$  levels using ANOVA between *APOE*  $\epsilon$ 4 carriers and noncarriers in a boxplot representation with outlier analysis. (PDF 93 kb)

**Additional file 8: Figure S2.** Scatter plots for FBB-PET global SUVR and A $\beta$  plasma ratios in SCD subjects. Correlations between plasma biomarkers and brain A $\beta$  burden. Biomarkers values plotted against SUVR values from FBB-PET imaging: FP42/40 (A), BP42/40 (B), FP42/TP42 (C), and FP40/TP40 (D). (PDF 286 kb)

**Additional file 9: Figure S3.** Linear regression between FBB-PET and A $\beta$  TP42/40 plasma ratio in *APOE*  $\epsilon$ 4 stratification SCD population. A) *APOE*  $\epsilon$ 4 carriers; B) *APOE*  $\epsilon$ 4 noncarriers. (PDF 115 kb)

**Additional file 10: Table S5.** Interaction between *APOE* and L<sub>TP42/40</sub>. (DOCX 14 kb)

**Additional file 11: Figure S5.** Box plots for TP40 and TP42 by FBB-PET global SUVR status in SCD subjects. (JPEG 53 kb)

## Abbreviations

ABtest: Araclon Biotech test; AD: Alzheimer's disease; ANOVA: Analysis of variance; *APOE*: Apolipoprotein E; *APP*: Amyloid precursor protein; AUROC: Area under the receiver operating characteristic curve; A $\beta$ : Amyloid beta; BP: Bound to plasma components; CDR: Clinical Dementia Rating; CI: Confidence interval; CSF: Cerebrospinal fluid; ELISA: Enzyme-linked immunosorbent assay; FACEHBI: Fundació ACE Healthy Brain Initiative; FBB: Florbetaben; FLAIR: Fast fluid-attenuated inversion recovery; FOV: Field of view; FP: Free plasma; FSL: FMRIB Software Library; HADS: Hospital Anxiety and Depression Scale; LLQ: Lower limit of quantification; MCI: Mild cognitive impairment; MFE: Memory Failures in Everyday Life Questionnaire; MMSE: Mini-Mental State Examination; MPRAGE: Magnetization-prepared rapid gradient-echo; MRI: Magnetic resonance imaging; NBACE: Fundació ACE Neuropsychological Battery; NPV: Negative predictive value; OR: Odds ratio; PET: Positron emission tomography; PPV: Positive predictive value; SCD: Subjective cognitive decline; SUVR: Standardized uptake value ratio; TE: Echo time; TI: Inversion time; TP: Total plasma; TR: Repetition time; ULQ: Upper limit of quantification

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## Availability of data and materials

The datasets used and/or analyzed during the present study are available from the corresponding author on reasonable request.

## Authors' contributions

AR, LT, MB, MS, and PP contributed to the study concept and design. IdR analyzed, interpreted and drafted the manuscript. AR and PP contributed to the literature search and drafted the manuscript. AR and SV contributed to analysis and interpretation of data. OR-G, AS, AP-C, CA, IH, MR-R, AM, LV, MA, AE, GO, MG, AG, MAS-S, SR, LM, EM, EP, AO, and SM-G were involved in data collection, recruitment and evaluation of the patients. JR, PP, VP-G, and MS participated in analytical data acquisition. FL, FC, AV, MG-C, MAT, and JG performed MRI and FBB-PET assessments. OS-G and GM-R analyzed the neuroimaging data. All authors read and approved the final manuscript.

## Ethics approval and consent to participate

The FACEHBI protocol received approval from the ethics review board of the Hospital Clínic i Provincial (Barcelona, Spain) (EudraCT: 2014-000798-38). All the participants signed written informed consent prior to any evaluation according to Spanish biomedical laws and to the principles of the Declaration of Helsinki.

## Consent for publication

Not applicable.

## Competing interests

JR, PP, VP-G, and MS are employees of Araclon Biotech Ltd. MS holds several patents related to Alzheimer's disease diagnosis and treatment, and he is the founder, chief executive officer, chief scientific officer, and one of the current shareholders of Araclon Biotech Ltd. The remaining authors declare that they have no competing interests.

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