

## PDF hosted at the Radboud Repository of the Radboud University Nijmegen

The following full text is a publisher's version.

For additional information about this publication click this link.

<http://hdl.handle.net/2066/137178>

Please be advised that this information was generated on 2017-12-05 and may be subject to change.



# Variability of CSF Alzheimer's Disease Biomarkers: Implications for Clinical Practice

Stephanie J. B. Vos<sup>1\*</sup>, Pieter Jelle Visser<sup>1,2</sup>, Frans Verhey<sup>1</sup>, Pauline Aalten<sup>1</sup>, Dirk Knol<sup>3</sup>, Inez Ramakers<sup>1</sup>, Philip Scheltens<sup>2</sup>, Marcel G. M. Olde Rikkert<sup>4,5</sup>, Marcel M. Verbeek<sup>5,6</sup>, Charlotte E. Teunissen<sup>7</sup>

**1** Department of Psychiatry and Neuropsychology, Maastricht University, School for Mental Health and Neuroscience, Alzheimer Center Limburg, Maastricht, the Netherlands, **2** Alzheimer center & Department of Neurology, Neuroscience Campus Amsterdam, VU University Medical Center, Amsterdam, The Netherlands, **3** Department of Epidemiology and Biostatistics, VU University Medical Center, Amsterdam, the Netherlands, **4** Department of Geriatric Medicine, Radboud University Nijmegen Medical Centre, Nijmegen, The Netherlands, **5** Donders Institute for Brain, Cognition and Behaviour, Radboud University Nijmegen and Radboud Alzheimer Centre, Nijmegen, The Netherlands, **6** Department of Neurology and Department of Laboratory Medicine, Radboud University Nijmegen Medical Centre, Nijmegen, The Netherlands, **7** Department of Clinical Chemistry, Neurological Laboratory, VU University Medical Center, Amsterdam, The Netherlands

## Abstract

**Background:** Cerebrospinal fluid (CSF) biomarkers are increasingly being used for diagnosis of Alzheimer's disease (AD).

**Objective:** We investigated the influence of CSF intralaboratory and interlaboratory variability on diagnostic CSF-based AD classification of subjects and identified causes of this variation.

**Methods:** We measured CSF amyloid- $\beta$  (A $\beta$ ) 1-42, total tau (t-tau), and phosphorylated tau (p-tau) by INNOTEST enzyme-linked-immunosorbent assays (ELISA) in a memory clinic population (n = 126). Samples were measured twice in a single or two laboratories that served as reference labs for CSF analyses in the Netherlands. Predefined cut-offs were used to classify CSF biomarkers as normal or abnormal/AD pattern.

**Results:** CSF intralaboratory variability was higher for A $\beta$ 1-42 than for t-tau and p-tau. Reanalysis led to a change in biomarker classification (normal vs. abnormal) of 26% of the subjects based on A $\beta$ 1-42, 10% based on t-tau, and 29% based on p-tau. The changes in absolute biomarker concentrations were paralleled by a similar change in levels of internal control samples between different assay lots. CSF interlaboratory variability was higher for p-tau than for A $\beta$ 1-42 and t-tau, and reanalysis led to a change in biomarker classification of 12% of the subjects based on A $\beta$ 1-42, 1% based on t-tau, and 22% based on p-tau.

**Conclusions:** Intralaboratory and interlaboratory CSF variability frequently led to change in diagnostic CSF-based AD classification for A $\beta$ 1-42 and p-tau. Lot-to-lot variation was a major cause of intralaboratory variability. This will have implications for the use of these biomarkers in clinical practice.

**Citation:** Vos SJB, Visser PJ, Verhey F, Aalten P, Knol D, et al. (2014) Variability of CSF Alzheimer's Disease Biomarkers: Implications for Clinical Practice. PLoS ONE 9(6): e100784. doi:10.1371/journal.pone.0100784

**Editor:** Gianluigi Zanusso, University of Verona, Italy

**Received:** April 17, 2014; **Accepted:** May 28, 2014; **Published:** June 24, 2014

**Copyright:** © 2014 Vos et al. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

**Data Availability:** The authors confirm that all data underlying the findings are fully available without restriction. All relevant data are within the paper and its Supporting Information files.

**Funding:** This study was funded by the Center for Translational Molecular Medicine (www.ctmm.nl) project LeARN (grant 02N-01) and in part by Zon-Mw as part of the BIOMARKAPD project of the European Joint Programming Initiative on Neurodegenerative Disorders (JPND). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

**Competing Interests:** Dr. Visser has served as an advisory board member of Bristol-Myers Squibb. He receives/received research grants from Zon-Mw as part of the BIOMARKAPD project in the frame of the European Joint Programming Initiative on Neurodegenerative Disorders (JPND), Innovative Medicines Initiative (IMI) as part of the EMIF-AD project, Bristol-Myers Squibb, European Commission 6th and 7th Framework programme, Life Sciences, Genomics and Biotechnology for Health. Dr. Verhey serves on the advisory board of Nutricia Medical Food bv. Dr. Scheltens serves/has served on the advisory boards of Genentech, Novartis, Roche, Danone, Nutricia, Baxter, and Lundbeck. He has been a speaker at symposia organized by Lundbeck, Merz, Danone, Novartis, Roche, and Genentech. For all his activities he receives no personal compensation. He is a member of the scientific advisory board of the EU Joint Programming Initiative and the French National Plan Alzheimer. The Alzheimer Center receives unrestricted funding from various sources through the VUmc Fonds. Dr. Teunissen serves at the advisory boards of Roche and Innogenetics. The authors confirm that this does not alter the authors' adherence to PLOS ONE policies on sharing data and materials.

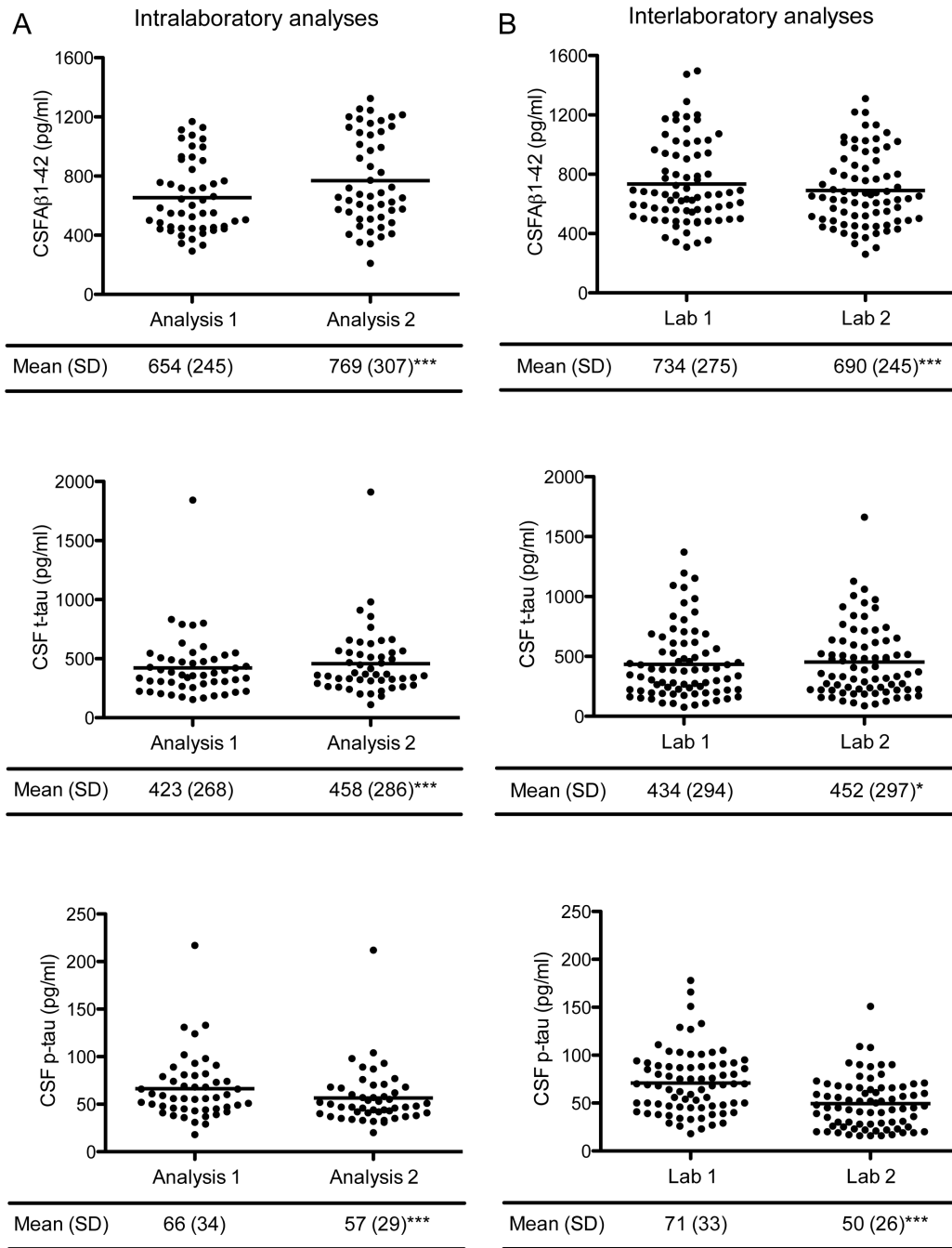
\* Email: s.vos@maastrichtuniversity.nl

## Introduction

Amyloid- $\beta$  (A $\beta$ ) 1-42, total tau (t-tau), and phosphorylated tau (p-tau) proteins in cerebrospinal fluid (CSF) are well-established biomarkers for Alzheimer's disease (AD) [1–3], and are increasingly being used for diagnosis in clinical practice. Previous studies reported considerable intra- or interlaboratory variability of CSF

analyses [4–7], which may influence the diagnostic classification. In this study, we performed a large-scale CSF multicenter study and investigated the exact influence of intra- and interlaboratory variability on CSF-based AD classification of subjects.

We hypothesized that the change of diagnosis would be largest for classification of subjects based on CSF A $\beta$ 1-42, as previous studies showed larger variability for CSF A $\beta$ 1-42 than for t-tau



**Figure 1. CSF levels by analysis and marker.** Results are frequencies and mean (SD) for each CSF marker on the left (A) for CSF intralaboratory analyses and on the right (B) for CSF interlaboratory analyses. The solid line represents the mean CSF levels. Analysis 1 is routine practice and analysis 2 is performed as part of the LeARN study. CSF = cerebrospinal fluid, Aβ = amyloid beta, t-tau = total tau, p-tau = phosphorylated tau. \*\*P<0.001, \*p<0.05 compared to CSF analysis 1 or lab 1. doi:10.1371/journal.pone.0100784.g001

and p-tau [5–7]. It was also hypothesized that change of diagnosis would be lower for analyses performed in the same laboratory than for analyses performed in different laboratories because CSF intralaboratory variability has been reported to be lower than interlaboratory variability (2.3–25% vs. 13–38%) [5–7].

We investigated intralaboratory and interlaboratory variability of CSF Aβ1-42, t-tau, and p-tau analyses by INNOTEST enzyme-linked-immunosorbent assays (ELISA). Samples were measured

twice in one or two laboratories that served as reference labs for AD CSF analyses in the Netherlands. We classified subjects based on validated cut-offs and examined how often CSF-based AD diagnosis changed after the second analysis.

**Table 1.** Agreement between CSF analyses.

	Intralaboratory analyses		Interlaboratory analyses	
	Correlation*	ICC*	Correlation*	ICC*
CSF A $\beta$ 1-42	0.85 (0.74–0.91)	0.76 (0.43–0.89)	0.94 (0.91–0.96)	0.92 (0.85–0.96)
CSF t-tau	0.98 (0.97–0.99)	0.97 (0.92–0.99)	0.98 (0.96–0.99)	0.98 (0.96–0.99)
CSF p-tau	0.95 (0.92–0.97)	0.90 (0.53–0.97)	0.94 (0.90–0.96)	0.73 (0.07–0.92)

Results are Pearson correlation and ICC (95% CI) for intra- and interlaboratory CSF analyses. ICC = Intraclass coefficients, ratio = A $\beta$ 1-42/t-tau, CSF = cerebrospinal fluid, A $\beta$  = amyloid beta, t-tau = total tau, p-tau = phosphorylated tau.

\*All  $p < 0.001$ .

doi:10.1371/journal.pone.0100784.t001

## Materials and Methods

### CSF Samples

CSF samples were collected from subjects included in the Leiden Alzheimer Research Netherlands (LeARN) study [8]. LeARN is a Dutch multicenter study performed in a memory-clinic setting that included subjects between October 2009 and May 2011 that had been newly referred for the assessment of cognitive complaints. Inclusion criteria were baseline diagnosis of subjective cognitive impairment (SCI), mild cognitive impairment (MCI) or dementia, Mini-Mental State Examination (MMSE)  $\geq$  20, clinical dementia rating scale (CDR) of maximal 1. Exclusion criteria were somatic, psychiatric or neurological disorders that could have caused the cognitive impairment.

In 3 of the 4 participating centers (Amsterdam, Maastricht, and Nijmegen), 126 CSF samples were analyzed in twofold, the first time as part of clinical routine and secondly for the LeARN study (Figure S1). For routine practice, samples were analyzed in Amsterdam and Nijmegen, which serve as national CSF AD biomarker centers. For the LeARN study, all samples were analyzed in a single batch in Amsterdam. The samples collected in Amsterdam ( $n = 50$ ) were measured twice in the same lab and were used to assess intralaboratory variability. Samples collected in Nijmegen ( $n = 32$ ) and Maastricht ( $n = 44$ ) were measured twice in different laboratories, i.e. in Nijmegen for clinical routine and in Amsterdam for the LeARN study, and were used to study interlaboratory variability (total  $n = 76$ ). Table S1 provides baseline patient demographics. The medical research ethics committee in Maastricht and the institutional review boards of Maastricht University Medical Center, VU University Medical Center Amsterdam, Radboud University Nijmegen Medical Center, and Leiden University Medical Center approved the study. All subjects provided written informed consent.

### CSF Procedures and Analyses

CSF was obtained by lumbar puncture between the L3/L4 or L4/L5 intervertebral space, and collected and aliquoted into polypropylene tubes. Samples for clinical routine of Maastricht were transported the same day on room temperature or stored at  $-20^{\circ}\text{C}$  and transported on dry ice within one week to the Nijmegen laboratory for analysis. Samples for clinical routine analysis were stored at  $-20^{\circ}\text{C}$  for up to 4 weeks (Amsterdam cohort) or at  $-80^{\circ}\text{C}$  for up to 2 weeks (Maastricht/Nijmegen cohort) before analysis. Research samples (i.e. samples of the LeARN study) were stored at  $-80^{\circ}\text{C}$  [9], at each center and samples of Maastricht and Nijmegen were transported on dry ice to Amsterdam for analysis after up to 2.5 years. Both laboratories used the commercially available INNOTEEST enzyme-linked

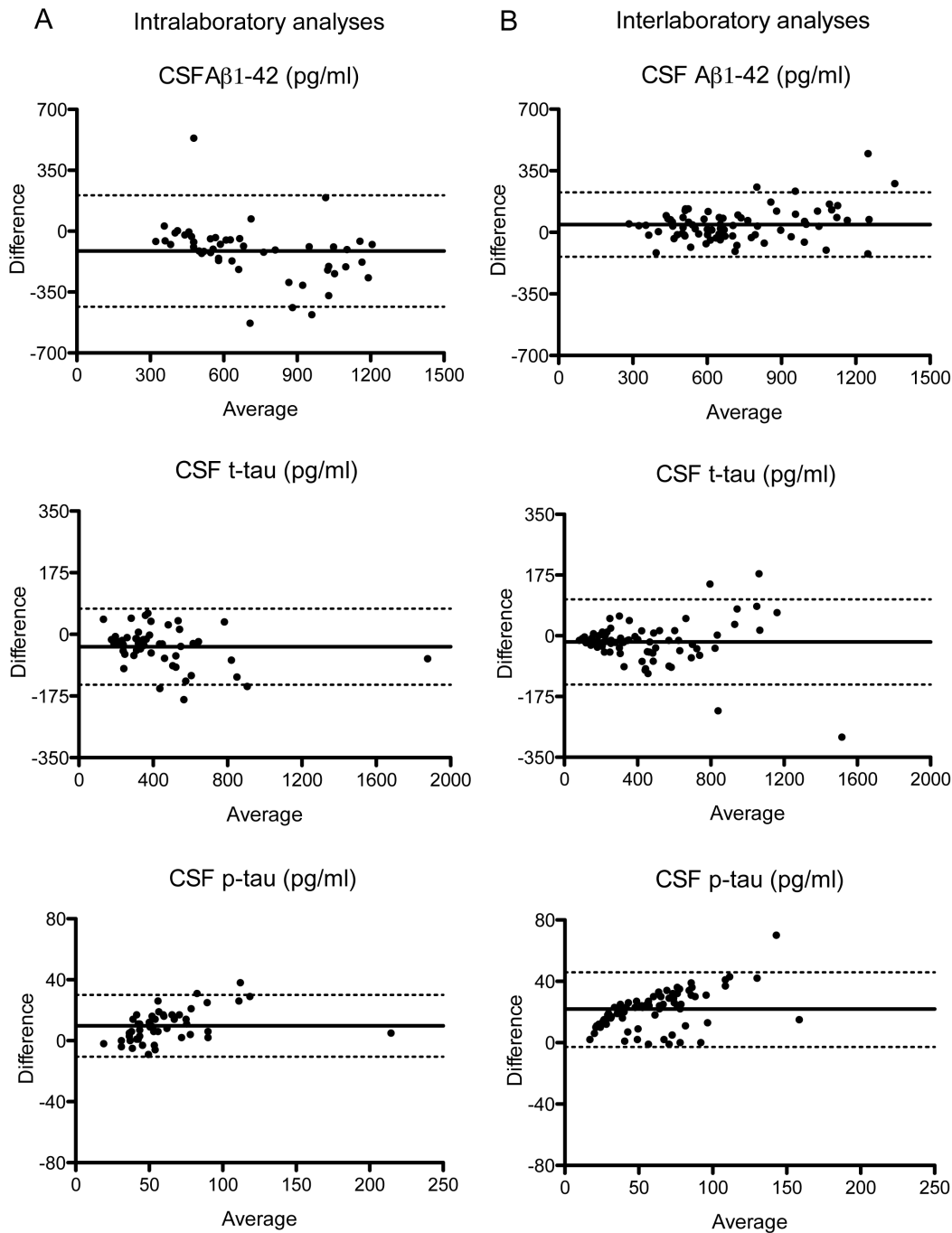
immunosorbent assays (ELISAs; Innogenetics, Ghent, Belgium) to quantify CSF A $\beta$ 1-42, CSF t-tau, and CSF p-tau, all performed by experienced laboratory technicians. For analysis of the research samples, the same lot number was used for all analyses, while for clinical routine analyses different lots were used in Nijmegen as well as in Amsterdam. Due to insufficient fluid material, CSF p-tau values were only available for 49 samples for intralaboratory analyses.

We also analyzed internal control samples from the Amsterdam lab to investigate the influence of lot-to-lot variation on measured CSF concentrations. One control sample had an AD typical profile and the other a normal CSF profile. The internal controls were aliquots for single use obtained by pooling surplus CSF. These aliquots were stored at  $-80^{\circ}\text{C}$  and all internal controls used in the current study were from the same batch of pools.

To study differences in biomarker classification as normal versus abnormal between CSF measurements, we dichotomized the CSF variables according to routinely used validated cut-offs of each lab. In Amsterdam, cut-offs were determined that could differentiate subjects with SCI from subjects with AD-type dementia with 85% sensitivity: CSF A $\beta$ 1-42  $\leq$  550 pg/ml, t-tau  $>$  375 pg/ml, and p-tau  $>$  52 pg/ml [10]. In Nijmegen, cut-offs were determined that could differentiate cognitively normal controls from subjects with AD-type dementia with a specificity of 95%: CSF A $\beta$ 1-42  $<$  500 pg/ml, t-tau  $>$  350 pg/ml, and p-tau  $>$  85 pg/ml [11]. Given that a different approach was used to define cut-offs in Amsterdam and Nijmegen, we performed interlaboratory analyses with the same cut-offs (Amsterdam cut-offs) as well as with lab-specific cut-offs (Amsterdam and Nijmegen cut-offs).

### Statistical Analyses

Statistical analyses were done with SPSS version 19.0 (Chicago, IL, USA) and GraphPad Prism 5, with significance set at  $p < 0.05$ . Intralaboratory and interlaboratory coefficients of variation (CV) were calculated as the standard deviation (SD) divided by the mean of the measurements of each sample for each biomarker. Subsequently, a mean CV was calculated. We performed paired t-tests to investigate the intralaboratory and interlaboratory variability between CSF analyses of CSF A $\beta$ 1-42, t-tau, and p-tau. In addition, we calculated Pearson correlations  $r$  and Intraclass Correlation Coefficients (ICC). An ICC score ranges from 0 to 1, representing virtually no (0.00–0.10), a slight (0.11–0.40), fair (0.41–0.60), moderate (0.61–0.80), or substantial (0.81–1.00) level of agreement between the analyses [12]. Confidence intervals were calculated for both the Pearson correlations and ICC scores. We made Bland-Altman plots to visualize the agreement between the CSF analyses [13], and calculated the percentage of subjects with change of AD marker classification



**Figure 2. Bland-Altman plots of variability between CSF analyses.** The average of CSF analysis 1 and 2 is plotted against the difference between both analyses, on the left (A) for CSF intralaboratory analyses and on the right (B) for CSF interlaboratory analyses. The solid line represents the mean and the dotted lines the upper and lower 1.95 SD. CSF = cerebrospinal fluid, A $\beta$  = amyloid beta, t-tau = total tau, p-tau = phosphorylated tau.

doi:10.1371/journal.pone.0100784.g002

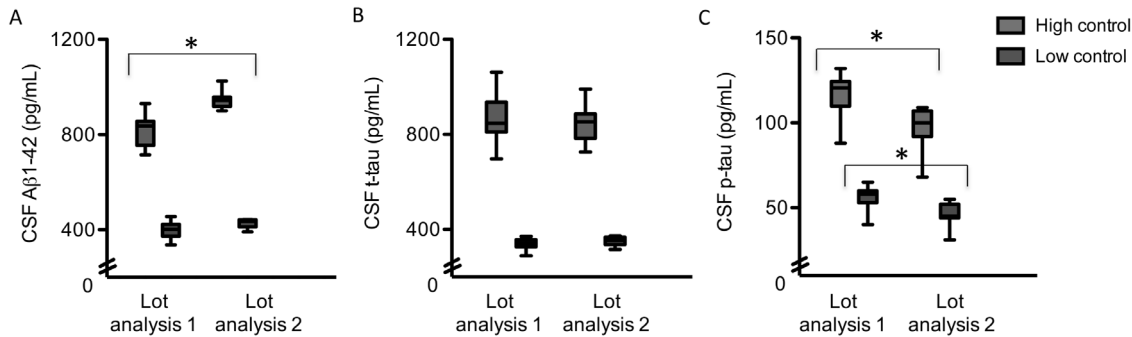
after the second analysis using routine CSF cut-offs of each laboratory. Interlaboratory lot-to-lot variation was examined using Deming regression [14].

## Results

### Intralaboratory Variability

CSF A $\beta$ 1-42 and t-tau levels were higher and p-tau levels were lower after reanalysis in the same laboratory ( $p < 0.05$  for all

analysis, Figure 1A, Figure S2A). The mean intralaboratory CV was 14.4% for A $\beta$ 1-42, 8.5% for t-tau, and 12.6% for p-tau. For CSF A $\beta$ 1-42, the correlation (0.85) and ICC (0.76) were moderate and lower than that of CSF t-tau ( $r = 0.98$  and ICC = 0.97) and p-tau ( $r = 0.95$  and ICC = 0.90; Table 1, Figure 2A). Internal quality control samples that were analyzed at each measurement also showed higher A $\beta$ 1-42 levels, slightly higher t-tau levels, and lower p-tau levels in the second measurement compared to the first measurement (Figure 3). For the clinical analyses 2 lots were used



**Figure 3. Internal control data of the Amsterdam laboratory.** Results are changes in high and low CSF biomarker levels for intralaboratory reanalysis in two lots (routine lot and LeARN lot) as part of internal control of data at the Amsterdam laboratory. A) Aβ1-42: Lot 1 used from February 2010 to February 2011 (n = 18) and lot 2 used from March to October 2012 (n = 17). B) T-tau: Lot 1 used from February 2010 to April 2011 (n = 24) and lot 2 used from March to June 2012 (n = 11). C) P-tau: Lot 1 used from February 2010 to April 2011 (n = 23) and lot 2 used from March to October 2012 (n = 11). CSF = cerebrospinal fluid, Aβ = amyloid beta, t-tau = total tau, p-tau = phosphorylated tau. \*P < 0.001 for differences between lot of analysis 1 (routine lot) and lot of analysis 2 (LeARN lot). doi:10.1371/journal.pone.0100784.g003

(lot 1 and lot 2), although the majority was measured with the same lot (lot 1). The reanalysis of all samples was performed with another lot (lot 3). Head-to-head comparison of lot 1 and lot 3 for the samples in our study showed that particularly the difference in Aβ1-42 levels between measurements could be explained by lot-to-lot variation although the mean t-tau levels also differed between lot 1 and 3 (Figure 4).

### Interlaboratory Variability

CSF Aβ1-42 levels were lower, t-tau levels higher, and p-tau levels lower after reanalysis in the second laboratory ( $p < 0.05$  for all analysis, Figure 1B, Figure S2B). The interlaboratory CV was 7.3% for Aβ1-42, 6.7% for t-tau, and 27.6% for p-tau. For CSF p-tau, the correlation (0.94) and the ICC (0.73) were high to moderate and lower than that of CSF Aβ1-42 ( $r = 0.94$  and ICC = 0.92) and t-tau ( $r = 0.98$  and ICC = 0.98; Table 1, Figure 2B).

### Change in AD Classification

We investigated how reanalysis changed the CSF AD classification based on individual CSF markers and based on the combination of CSF markers with an AD profile being defined as abnormal Aβ1-42 and abnormal t-tau or p-tau. Using predefined cut-offs, repeated CSF analyses in the same laboratory led to a change in biomarker classification (normal vs. abnormal) of 26% of subjects based on Aβ1-42, 10% based on t-tau, 29% based on p-tau, and 16% based on the AD profile (Table 2). Repeated CSF analyses in different laboratories using the cut-offs from the Amsterdam lab led to a change in biomarker classification of 12% of subjects based on CSF Aβ1-42, 1% based on t-tau, 22% based on p-tau, and 14% based on the AD profile (Table 2). When we applied lab-specific cut-offs to define an abnormal score, the repeated CSF analyses in different laboratories led to a change in biomarker classification of 17% of subjects based on CSF Aβ1-42, 1% based on t-tau, 12% based on p-tau, and 12% based on the AD profile (Table 2). Figure 5 shows the change in CSF levels for each biomarker for intra- and interlaboratory analyses of subjects in whom reanalysis led to a different biomarker classification as normal vs. abnormal when Amsterdam cut-offs were applied. While most of the subjects with a change in AD classification after reanalysis in the same laboratory as well as in a different laboratory had CSF biomarker values relatively close to the cut-off points, some showed larger changes in CSF biomarker values.

The mean change in CSF values for intralaboratory analyses was 177 pg/ml (95% CI 78–275) for Aβ1-42, 72 (13–130) for t-tau, and 12 (9–16) for p-tau. For interlaboratory analyses with the Amsterdam cut-offs, the mean change in CSF biomarker values was 153 pg/ml (57–250) for Aβ1-42 and 26 (20–32) for p-tau. For t-tau, only one subjects showed a change in biochemical diagnosis (a change of 44 pg/ml).

### Discussion

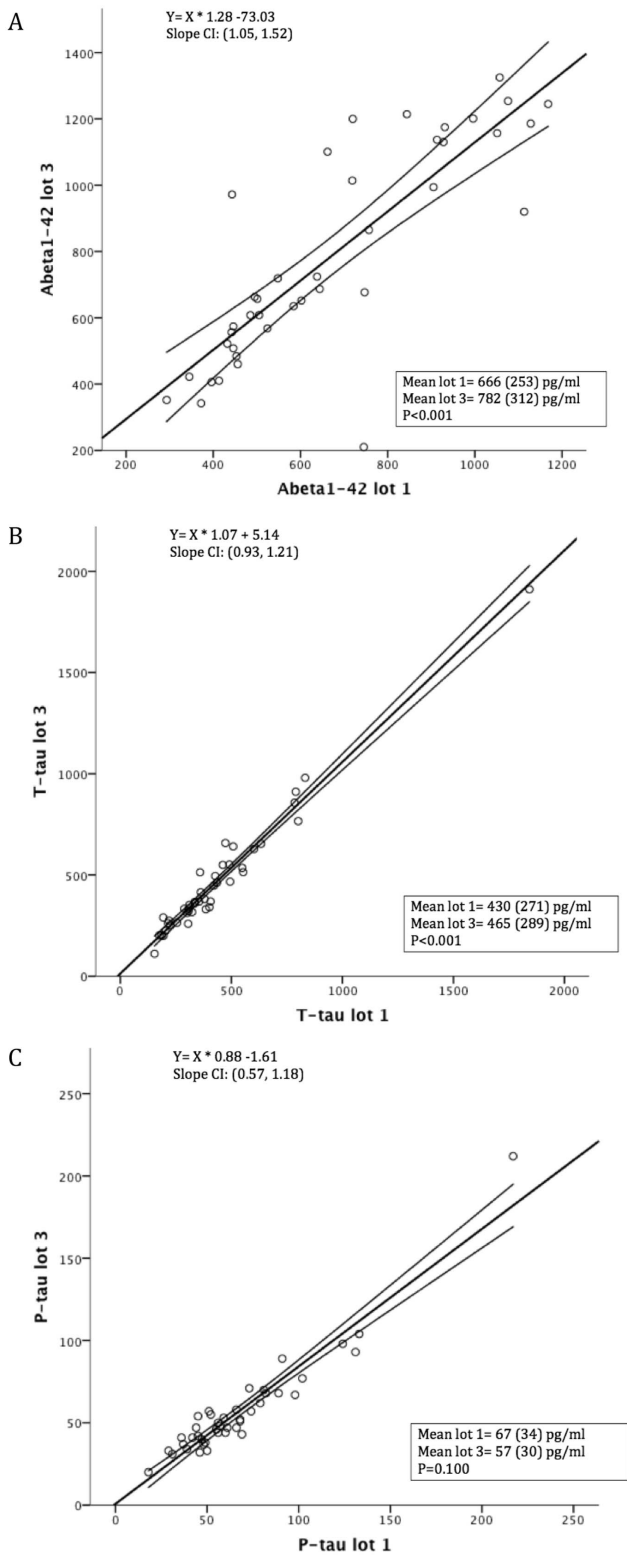
We observed clear variability in CSF AD biomarker levels between repeated analyses in the same laboratory as well as between two different laboratories. Our study is the first to show that this variability frequently led to a change in CSF-based AD diagnosis when predefined cut-offs for abnormal CSF values were applied.

CSF intralaboratory variability (based on CV) was largest for Aβ1-42, consistent with previous studies [5–7]. Change in AD-like scores after repeated analyses in the same laboratory was, however, highest for CSF p-tau (29%) followed by Aβ1-42 (26%) and t-tau (10%). This large change for p-tau is likely due to a smaller range of values of p-tau and values being closer to the cut-off compared to other markers, such that a small change in observed concentration more easily leads to a different classification.

CSF interlaboratory variability (based on CV) was largest for p-tau, unlike findings from previous studies, which found that interlaboratory variability was largest for Aβ1-42 [5–7]. Change in AD-like scores after repeated analyses in different laboratories was also higher for CSF p-tau (22%) than for Aβ1-42 (12%) and t-tau (1%) when the same cut-offs were used for each site. A rather unexpected finding was that the interlaboratory variability of Aβ1-42 was smaller than the intralaboratory variability.

A major finding was that CSF analytical variability frequently led to a change in diagnostic CSF-based AD classification. Importantly, also a diagnosis of a CSF AD profile based on Aβ1-42 and t-tau or p-tau changed in 12–16% of the cases. As some of the subjects whose AD classification changed after reanalysis had values around the cut-off points, it could be helpful to use a range around a cut-off point rather than a fixed cut-off point.

The intra- and interlaboratory variability in CSF results can result from differences in preanalytical and analytical procedures, and lot-to-lot variation of analytical kits [9,15]. The intralabora-



**Figure 4. Intralaboratory lot-to-lot variation.** Results in this graph are based on deming regression and show the CSF levels of lot 1 and lot 3 for Aβ1-42 (A), t-tau (B), and p-tau (C). The slope is different for Aβ1-42 levels between lot 1 and lot 3. The mean difference in CSF levels between lot 1 and 3 was significantly different for Aβ1-42 and t-tau. Lot 1 = clinical routine lot, lot 3 = LeARN lot, Aβ = amyloid beta, t-tau = total tau, p-tau = phosphorylated tau.  
 doi:10.1371/journal.pone.0100784.g004

**Table 2. Numbers of subjects with normal and abnormal classified CSF marker values based on predefined cut-offs.**

Classification of biomarker(s)	Intralaboratory analyses				Interlaboratory analyses				
	Amsterdam cut-offs		Amsterdam cut-offs*		Amsterdam cut-offs**		Amsterdam cut-offs**		
	Aβ1-42	P-tau	Aβ1-42	P-tau	Aβ1-42	P-tau	Aβ1-42	P-tau	
Analysis 1 & 2 same classification	74% (37)	90% (45)	71% (35)	84% (42)	88% (67)	78% (59)	86% (65)	83% (63)	88% (67)
Analysis 1 normal & 2 abnormal	2% (1)	4% (2)	6% (3)	-	11% (8)	-	9% (7)	16% (12)	11% (8)
Analysis 1 abnormal & 2 normal	24% (12)	6% (3)	23% (11)	16% (8)	1% (1)	22% (17)	5% (4)	1% (1)	12% (9)
					1% (1)	1% (1)	1% (1)	1% (1)	1% (1)

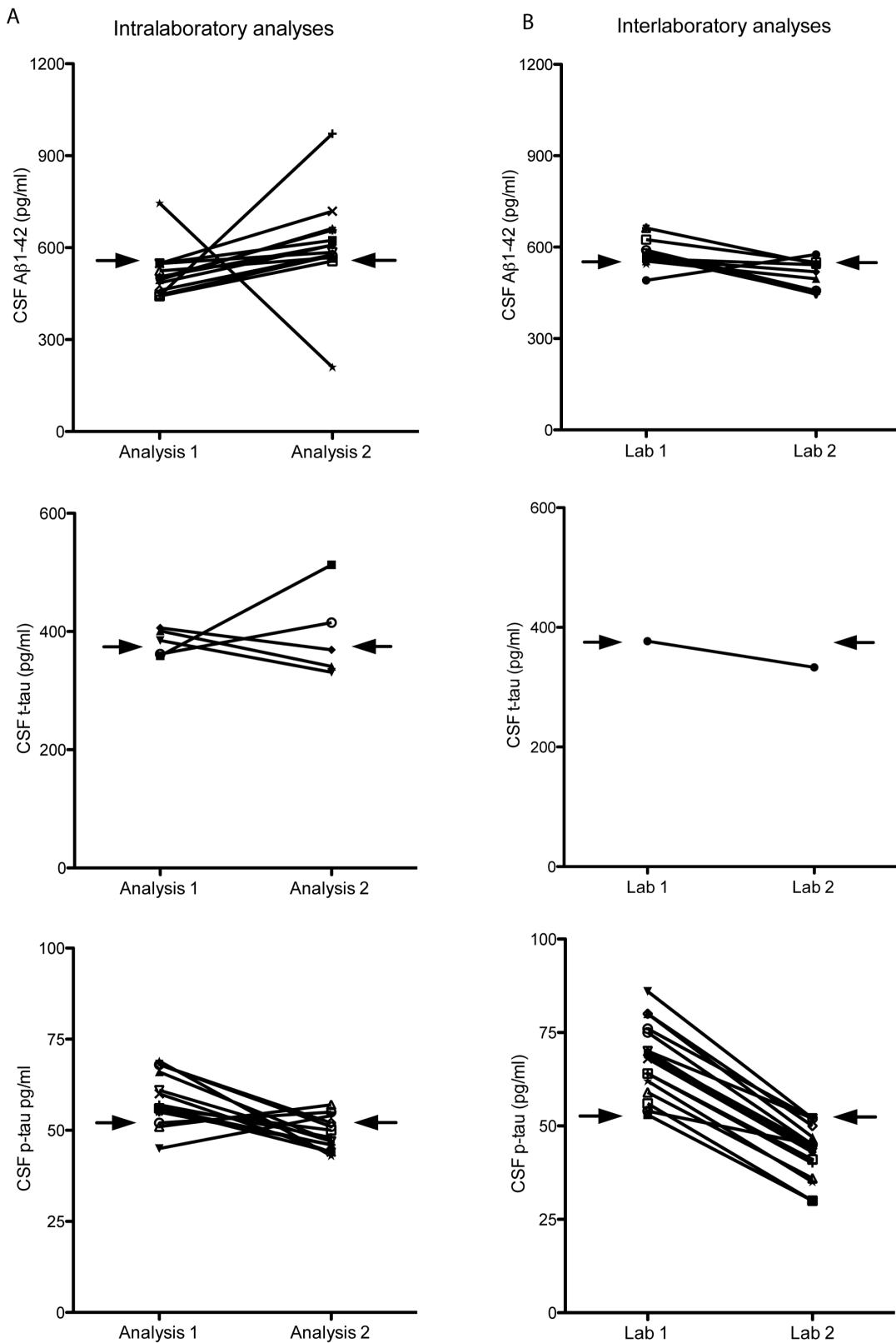
Results are the percentage (number) of subjects with normal and abnormal classified CSF markers based on CSF analysis 1 (routine practice) and analysis 2 (LeARN study). The CSF AD profile was defined as abnormal Aβ1-42 and abnormal t-tau or p-tau, based on predefined cut-offs (see methods).

CSF = cerebrospinal fluid, Aβ = amyloid beta, t-tau = total tau, p-tau = phosphorylated tau.

\*The routine analyses in Nijmegen and the LeARN analyses were dichotomized according to the Amsterdam cut-offs (see methods).

\*\*The routine analyses in Nijmegen were dichotomized according to the Nijmegen cut-offs and the LeARN analyses according to the Amsterdam cut-offs (see methods).

doi:10.1371/journal.pone.0100784.t002



**Figure 5. Change in CSF marker classification according to cut-offs.** Results are CSF levels only for subjects in which reanalysis let to a different biomarker classification using the cut-offs from Amsterdam to define abnormal CSF values. On the left (A) are changes in biomarker levels for intralaboratory reanalysis and on the right (B) for interlaboratory reanalysis. The arrow represents the applied Amsterdam CSF cut-off. Analysis 1 is routine practice and analysis 2 is performed as part of the LeARN study. CSF=cerebrospinal fluid, Aβ=amyloid beta, t-tau=total tau, p-tau=phosphorylated tau.  
doi:10.1371/journal.pone.0100784.g005



tory variability is likely mainly due to lot-to-lot variation as other procedures remained essentially the same. Lot-to-lot variation is a technical limitation of the ELISA method, which should also be considered when defining cut-offs and interpreting CSF values close to the cut-off. Additional support for this comes from our observation that internal control values showed a striking lot-to-lot variation of up to 20%. The change in internal control values was of the same order of magnitude and same direction of change as the change in CSF scores from the patient samples. Head-to-head comparison of lots for the samples in our study showed that mainly CSF A $\beta$ 1-42 variability could be explained by lot-to-lot variation. This likely explains why CSF intralaboratory variability was largest for A $\beta$ 1-42. Another possible explanation for the intralaboratory variability is variability in freezing conditions and storage time. However, previous studies showed that these factors have a minor impact on variability of CSF values [16,17].

Interlaboratory variability may also be caused by lot-to-lot variation. Indeed, lots used in each lab showed only minor overlap (data not shown). Differences in analytical procedures may also have contributed to the interlaboratory variability. However, both laboratories used similar protocols, were both trained in a hands-on workshop [9], and were similarly experienced.

Another source of variability between laboratories, which may influence CSF-based AD classification, is the difference in cut-offs used. We, therefore, tested interlaboratory variability for CSF-based AD classification both with the same CSF cut-offs as well as lab-specific cut-offs. While lab-specific cut-offs for A $\beta$ 1-42 and t-tau did not differ much, there was a large difference for p-tau (85 pg/ml in Nijmegen vs. 52 pg/ml in Amsterdam). This was also reflected in the interlaboratory variability in CSF-based AD classification. Using the same cut-off, abnormal p-tau was more common when samples were analyzed in Nijmegen than in Amsterdam. Change in classification was often due to subjects with scores around the cut-off (Figure 5B). Using the lab-specific cut-off, however, abnormal p-tau was more common when samples were analyzed in Amsterdam than in Nijmegen. Here change in classification was mainly due to differences in cut-offs. This clearly indicates that lab-specific cut-offs may also influence comparability between laboratories.

Our study has several limitations. The cut-offs that were used may have influenced our findings on change in CSF-based AD diagnosis. However, as no universal CSF cut-offs are available, we applied routinely used validated cut-offs.

One of the major strengths of this study was the large number of CSF samples used to study interlaboratory variability, as most previous studies were based only on a few samples [5–7]. Furthermore, our study is the first to show head-to-head comparison of lots within one center to directly address the issue of lot-to-lot variability. Therefore, our findings may provide a valuable addition to the described findings of the Alzheimer Association Quality Control program. In addition, our study design allows generalization to other CSF centers that analyze

CSF AD biomarkers using ELISA, as it reflects CSF procedures in general clinical practice.

Together, our findings suggest that variability in CSF analyses is common between and within laboratories, in particular for A $\beta$ 1-42 and p-tau. A substantial part of this variability seems to be explained by lot-to-lot variation of analytical kits. The variability has a large impact on CSF-based AD diagnosis or treatment decisions in clinical settings, suggesting that we should be careful when interpreting CSF findings and always interpret them within a clinical context [18], and with reference to internal standards. Also the use of age-adjusted cut-offs may be helpful, as tau levels are known to increase with age.

For the moment this rightly restricts the indication for CSF biomarker testing in diagnostic guidelines as complimentary and non-obligatory [19,20]. The recent consensus on standardization of preanalytical aspects of CSF analyses [15,17], as well as the ongoing worldwide quality control study [6], and standardization projects ([www.neurodegenerationresearch.eu](http://www.neurodegenerationresearch.eu)) will help to move towards a standardized and harmonized implementation of CSF markers in clinical routine.

## Supporting Information

**Figure S1 Laboratories that performed CSF analyses for clinical routine are presented on the left (analysis 1); the laboratory that performed CSF analyses for the LeARN study is presented on the right (analysis 2).** (DOCX)

**Figure S2 Change in CSF levels after reanalysis.** Results on the left (A) are changes in CSF biomarker levels for intralaboratory reanalysis and on the right (B) for interlaboratory reanalysis. Analysis 1 is routine practice and analysis 2 is performed as part of the LeARN study. CSF = cerebrospinal fluid, A $\beta$  = amyloid beta, t-tau = total tau, p-tau = phosphorylated tau. (DOCX)

**Table S1 Baseline patient demographics.** Results are mean (SD) or number (%), presented for the total sample and separately for the sample for intralaboratory analyses (Amsterdam cohort) and the sample for interlaboratory analyses (Maastricht & Nijmegen cohort). MMSE = Mini-Mental State Examination, CDR = Clinical dementia rating scale, SCI = subjective cognitive impairment, MCI = mild cognitive impairment, AD = Alzheimer's disease. (DOCX)

## Author Contributions

Conceived and designed the experiments: SV FV. Performed the experiments: MV CT. Analyzed the data: SV DK. Contributed to the writing of the manuscript: SV PJV CT. Reviewed the manuscript for intellectual content and approved the final draft: SV PJV FV PA DK IR PS MO MV CT. Involved in data collection: SV FV PA IR PS MO.

## References

- Blennow K, Hampel H (2003) CSF markers for incipient Alzheimer's disease. *Lancet Neurol* 2: 605–613.
- Fagan AM, Holtzman DM (2010) Cerebrospinal fluid biomarkers of Alzheimer's disease. *Biomark Med* 4: 51–63.
- Vos SJB, van Rossum IA, Verhey F, Knol DL, Soininen H, et al. (2013) Prediction of Alzheimer's disease in subjects with amnesic and non-amnesic MCI. *Neurology* 80: 1124–1132.
- Dumurgier J, Vercurysse O, Paquet C, Bombois S, Chaulet C, et al. (2012) Intersite variability of CSF Alzheimer's disease biomarkers in clinical setting. *Alzheimers Dement*. DOI:10.1016/j.jalz.2012.06.006
- Lewczuk P, Beck G, Ganslandt O, Esselmann H, Deisenhammer F, et al. (2006) International quality control survey of neurochemical dementia diagnostics. *Neurosci Lett* 409: 1–4.
- Mattsson N, Andreasson U, Persson S, Arai H, Batish SD, et al. (2011) The Alzheimer's Association external quality control program for cerebrospinal fluid biomarkers. *Alzheimers Dement* 7: 386–395.
- Verwey NA, van der Flier WM, Blennow K, Clark C, Sokolow S, et al. (2009) A worldwide multicentre comparison of assays for cerebrospinal fluid biomarkers in Alzheimer's disease. *Ann Clin Biochem* 46: 235–240.
- Handels RL, Aalten P, Wolfs CA, Olde Rikkert M, Scheltens P, et al. (2012) Diagnostic and economic evaluation of new biomarkers for Alzheimer's disease: the research protocol of a prospective cohort study. *BMC Neurol* 12: 72.

9. Teunissen CE, Verwey NA, Kester MI, van Uffelen K, Blankenstein MA (2010) Standardization of Assay Procedures for Analysis of the CSF Biomarkers Amyloid beta(1-42), Tau, and Phosphorylated Tau in Alzheimer's Disease: Report of an International Workshop. *Int J Alzheimers Dis*. doi:10.4061/2010/635053.
10. Mulder C, Verwey NA, van der Flier WM, Bouwman FH, Kok A, et al. (2010) Amyloid-beta(1-42), total tau, and phosphorylated tau as cerebrospinal fluid biomarkers for the diagnosis of Alzheimer disease. *Clin Chem* 56: 248–253.
11. de Jong D, Kremer BP, Olde Rikkert MG, Verbeek MM (2007) Current state and future directions of neurochemical biomarkers for Alzheimer's disease. *Clin Chem Lab Med* 45: 1421–1434.
12. Shrout PE (1998) Measurement reliability and agreement in psychiatry. In *Stat Methods Med Res*, 7: 301–317.
13. Bland JM, Altman DG (1986) Statistical methods for assessing agreement between two methods of clinical measurement. *Lancet* 1: 307–310.
14. Dunn G (2007) Regression models for method comparison data. *Journal of Biopharmaceutical Statistics* 17: 739–756.
15. del Campo M, Mollenhauer B, Bertolotto A, Engelborghs S, Hampel H, et al. (2012) Recommendations to standardize preanalytical confounding factors in Alzheimer's and Parkinson's disease cerebrospinal fluid biomarkers: an update. *Biomark Med* 6: 419–430.
16. Schoonenboom NS, Mulder C, Vanderstichele H, Van Elk EJ, Kok A, et al. (2005) Effects of processing and storage conditions on amyloid beta(1-42) and tau concentrations in cerebrospinal fluid: implications for use in clinical practice. *Clin Chem* 51: 189–195.
17. Vanderstichele H, Bibl M, Engelborghs S, Le Bastard N, Lewczuk P, et al. (2012) Standardization of preanalytical aspects of cerebrospinal fluid biomarker testing for Alzheimer's disease diagnosis: a consensus paper from the Alzheimer's Biomarkers Standardization Initiative. *Alzheimers Dement* 8: 65–73.
18. Spies PE, Claassen JA, Peer PG, Blankenstein MA, Teunissen CE, et al. (2012) A prediction model to calculate probability of Alzheimer's disease using cerebrospinal fluid biomarkers. *Alzheimers Dement*; doi:10.1016/j.jalz.2012.01.010.
19. Hort J, O'Brien JT, Gainotti G, Pirttila T, Popescu BO, et al. (2010). EFNS guidelines for the diagnosis and management of Alzheimer's disease. *Eur J Neurol* 17: 1236–1248.
20. Jack CR, Albert MS, Knopman DS, McKhann GM, Sperling RA, et al. (2011) Introduction to the recommendations from the National Institute on Aging-Alzheimer's Association workgroups on diagnostic guidelines for Alzheimer's disease. *Alzheimers Dement* 7: 257–262.