

AN UPDATE ON FLUID BIOMARKERS FOR NEURODEGENERATIVE DISEASES: RECENT SUCCESS AND CHALLENGES AHEAD.

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ABSTRACT

Over the last twenty years, the characterization of Alzheimer's disease (AD) patients has progressed from a description of clinical symptomatology followed by neuropathological findings at autopsy to detailed *in vivo* pathophysiological signatures using cerebrospinal fluid (CSF) and positron emission tomography (PET). Additionally, CSF biomarkers now also reflect synaptic pathology, axonal injury and neuroinflammation. Technological advances have provided ultrasensitive techniques capable of measuring proteins of pathophysiological importance at femtomolar concentrations in blood samples (*e.g.* amyloid, tau species and neurofilaments). This has huge potential to screen large populations in the near future, which is essential for secondary prevention trials and primary care management. For most neurodegenerative diseases, however, research has not reached the same success. Common pathologies, such as that underlying dementia with Lewy bodies, Parkinson's disease and frontotemporal dementias, are still without reliable diagnostic biomarkers, although emerging techniques show promising pilot results for some of these diseases. This is likely to change in the next few years, which will be of great importance to stratify populations enrolling in clinical trials, given that these pathologies often coexist.

INTRODUCTION

Neurodegenerative diseases (NDD) comprise a group of neurological conditions that share the trait of nerve cell death, together with signature protein inclusions found in autopsies, which is why they are sometimes referred to as proteinopathies. NDDs include a large number of diseases, but most can be classified by the presence of amyloid-beta ($A\beta$), tau, α -synuclein, prion protein (PrP), and transactive response DNA-binding protein 43 (TDP-43), resulting in cerebral amyloidosis, tauopathies, α -synucleinopathies, prion diseases, TDP-43 proteinopathies, respectively. Neuropathological examinations have demonstrated that these protein signatures often coexist, which is seen at a rate of 27-68% among the groups examined, according to several recent studies [1-3]. Among NDDs, Alzheimer's disease (AD) is the most common, and is the leading cause of dementia worldwide, representing 50-75% of the cases [4]. It is also the disease in which the search for reliable fluid biomarkers has been the most successful, with the combination of decreased $A\beta_{42}$ (particularly when examined in a ratio with $A\beta_{40}$), and increased tau phosphorylated at threonine residue 181 (p-tau181) and T-tau in cerebrospinal fluid (CSF) constituting the typical CSF biomarker signature in AD, which is now also included in the research criteria for AD [5]. However, because of the large majority of failed disappointing results from all clinical trials attempting to target AD progression, most clinical

trials now attempt to target AD and other NDDs at a preclinical stage as well as more accurately characterizing the individuals included. One example highlighting the need of a reliable and scalable biological characterization of pathology is the recent publication of a paper by a consensus work group describing a recently discovered disease entity mimicking AD, limbic-predominant age-related transactive domain binding protein 43 (TDP-43) encephalopathy (LATE), which is present in >20 % of autopsies conducted in individuals > 80 years old [6]. Therefore, the focus of much of current neurological biomarker research is to find reliable diagnostic and predictive biomarkers in more easily accessible biofluids, such as blood, at an early disease stage. Fortunately, research findings in recent years have generated promising results, which might soon be adopted in clinical and research settings, and will thus be the focus of this review together with recent progress in the field of CSF biomarkers. A summary of the biomarkers discussed can be found in figure 1. Since neurofilaments will be covered in other parts of this special issue, it will not be in the scope of this review.

FLUID BIOMARKERS FOR AMYLOID PATHOLOGY

Extracellular deposition of A β into plaques, through the cleavage of APP with β -secretase 1 (BACE1) and γ -secretase is proposed as the main pathogenic event in AD [7]. Since the amyloid cascade hypothesis was presented three decades ago, it has resulted in increased understanding of AD pathogenesis, as well as in reliable diagnostic biomarkers in CSF and through amyloid positron emission tomography (PET) imaging, reflecting cerebral amyloid burden [8].

CSF

In patients with autosomal dominant AD from the Dominantly Inherited Alzheimer's Network (DIAN), longitudinal data suggests that low CSF A β 42 compared to non-carriers precedes expected symptomatic AD with 25 years, followed by abnormalities in brain glucose metabolism a decade later, further supporting the role of A β deposition as the earliest sign and initiator of disease pathology [9]. As previously mentioned, even though A β 42 in CSF is an widely accepted diagnostic test for A β pathology in research criteria [5], as well as in the proposed biological staging of AD [10], with reduced CSF levels reflecting retention of A β 42 in the brain parenchyma, research is still ongoing in terms of standardizing protocols for CSF analyses in clinical laboratory practice [11]. The field is now moving towards using A β 42/40 or 38 ratios, due to their ability to correct for inter-individual differences in amyloid processing and possible preanalytical confounders [12]. CSF A β 42 and A β 42/40 are highly concordant with A β PET uptake [13]. However, in many cases, CSF A β 42 seemingly precedes A β PET

positivity [14] which reflects the ability of CSF to capture changes in oligomeric A β . Furthermore, in concordance with neuropathological findings that proteinopathies often coexist, A β 42 is decreased in most DLB patients [15] and many Parkinson disease (PD) dementia patients, but is normal in, *e.g.*, frontotemporal dementia (FTD) patients [16].

Blood

In contrast to earlier reports [17], recent findings suggest that plasma A β 42, especially in ratio with A β 40, reflect cerebral A β pathology. Current research demonstrates a good but not optimal correlation of plasma A β with cerebral A β pathology, as measured by PET or CSF. Promising results have been presented by several groups with immunoprecipitation mass spectrometry (IPMS) analyses [18] [19], as well as with immunoassays, such as Single molecule array (Simoa) [20, 21], enzyme-linked immunosorbent assay (ELISA) [22], and immunomagnetic reduction (IMR) assay [23, 24] (see Table 1). A recent validation study utilizing a fully automated immunoassay (Elecsys) to measure plasma A β 42 and A β 40 indicates the promising capabilities of plasma A β in clinical laboratory practice [25]. However, the concordance between studies is poor, but slightly better when comparing mass spectrometric assays [18, 19]. Immunoassays seemingly struggle with matrix effects possibly confounding the results, which are more effectively eliminated with IPMS techniques. This is reflected in the area under the receiver operation characteristics curve (AUC) ranging from 0.789-0.914. Schindler *et al.* report a difference between A β -positive and A β -negative of only 11 percent [19], this is compared to changes of around 50% in CSF [26]. Small effect sizes between A β -positive and A β -negative individuals, possibly due to peripheral A β expression [27], is an issue that needs to be addressed, if A β is to be employed as a screening test for cerebral amyloidosis. To elucidate these issues, cooperative efforts such as round robins and validation studies are warranted. Furthermore, studies have also explored the possibility of measuring plasma A β in heterogeneous forms [28] and exosomes [29], but data is still limited.

TAU PATHOLOGY

The aggregation of hyperphosphorylated forms of the axonal protein tau in the neuronal soma, forming neurofibrillary tangles (NFT), is a key neuropathological feature in AD, but tau inclusions in neurons or glial cells are also found in other NDDs [30].

CSF

The cornerstone markers T-tau and p-tau181 together with CSF A β 42 are proposed as biomarkers that biologically define AD [10]. They are considered diagnostic in the research criteria for AD [5]. T-tau is not different from HCs in most NDDs, except for AD and CJD [31]. However, CSF p-tau181 is seemingly AD-specific [32]. T-tau and p-tau181 correlates with tau-PET in later stages of the disease, but seemingly not in cognitively unimpaired AD patients, indicating that CSF biomarkers may be used as diagnostic markers, but not reflecting disease stage [33]. However, other tau fragments have been investigated as biomarkers reflecting tangle pathology. Increased baseline levels of the N-terminal fragment cleaved at residue 224 (N-224) distinguished AD from HC and patients with primary tauopathies, and predicted cognitive decline as well as conversion from MCI to AD [34]. A recent pilot study investigated a novel assay measuring C-terminally truncated (tau-368) tau found that lower levels of tau-368 was inversely correlated with disease stage, possibly indicating a decrease in tau production or more intriguingly sequestering of tau in NFTs [35]. It is important to note that this is a pilot study, meaning that these theories have to be further investigated. Additionally, recent data suggest that a novel assay targeting tau phosphorylated at 217 (p-tau217) very closely resembles the time of change and trajectory of p-tau181 in A β -positive individuals [36]. It is still not elucidated whether the increased p-tau181 and T-tau concentrations in CSF reflect tangle formation or neurodegeneration *per se*, as proposed in the research framework [10]. One study suggests that the increased CSF levels of tau are instead due to increased phosphorylation and secretion of tau in the CNS in response to brain amyloidosis [37]. Further, data from DIAN suggests that the increases of p-tau181 become less prominent in the symptomatic stages of the disease, and that this potentially reflects sequestering of phosphorylated tau in tangles [9]. This will be elucidated in further studies.

Blood

As previously stated, current research puts great effort in developing reliable screening and prognostic assays in blood for tau in NDDs, which has proven to be a difficult hurdle to overcome. These difficulties can partly be explained by the fact that tau is rapidly cleared from plasma (the apparent half-life of tau in blood is <24 hours), supported by studies conducted on patients with acute brain injury following cardiac arrest [38]. This is consistent with the findings that plasma T-tau correlates poorly with CSF T-tau [39]. Nevertheless, cross-sectional studies during the last 5 years have, in a fairly consistent manner, demonstrated that plasma T-tau levels

are increased in patients with AD according to a meta-analysis [17], but do not differentiate significantly between MCI and HC [40, 41]. Studies have shown that plasma T-tau is associated with cognitive decline as well as risk of MCI [41, 42] and AD [43], and that it correlates negatively with grey matter density [44], but this remains to be examined in longitudinal studies. However, the overlap seems to be large across diagnostic groups, indicating that plasma T-tau alone is not suitable as a diagnostic marker for AD. As previously mentioned, tau pathology is also present in other NDDs and higher plasma levels of T-tau have been observed in FTD [45, 46], PD, vascular dementia (VaD) and Creutzfeldt-Jakob disease (CJD) [46], where levels are substantially increased compared with HCs and correlate strongly with rate of disease progression [47, 48]. Targeting N-terminal fragments also in plasma might be a way forward, with one study separating AD and MCI from HCs with great sensitivity and specificity albeit small sample sizes [49]. The recent reports of detection of p-tau181 in plasma has generated much interest, with three studies being able to distinguish AD from HCs, using Single molecule array (Simoa) [50], immunomagnetic reduction (IMR) [51] and Meso Scale Discovery (MSD) [52] technologies, respectively. While IMR data suggests increases in most NDDs, MSD studies clearly show specific increases only in the AD continuum corroborating CSF findings. Mielke *et al.* originally demonstrated a correlation between p-tau181, and amyloid/tau PET, which indicates that plasma p-tau181 is a good predictor of brain AD pathology. These findings were replicated in a recent study by Palmqvist *et al.*, demonstrating that plasma p-tau181 associates with both A β PET positivity as well as CSF p-tau181. Interestingly, the change in plasma p-tau181 became significant before amyloid PET, but after CSF/plasma A β 42 [36]. Thus, plasma p-tau181 might be useful both diagnostically as well as for disease staging. Nonetheless, validation studies in larger cohorts, as well as with other emerging assays, are warranted to determine the clinical utility of plasma p-tau181 as an AD marker for primary care settings, specialized centers and clinic trials.

SYNAPTIC DEGENERATION

Synaptic degeneration is an early event in AD and other NDDs [53], and synaptic loss is the best correlate of cognitive decline [54]. Abnormal magnetic resonance imaging (MRI) or fluorodeoxyglucose PET scans are currently proposed as markers of synaptic loss in the proposed staging of AD [10]. Another emerging method to detect synaptic loss in NDDs is PET tracers targeting synaptic vesicle protein 2A (SV2A) [55]. In recent years, promising assays in biofluids targeting loss of synapses have been developed.

CSF

Neurogranin is a postsynaptic protein involved in long-term potentiation and synaptic plasticity and increased concentrations in CSF in AD patients accurately predict cognitive decline from MCI to AD dementia [56]. These increases are neuropathologically associated with plaque pathology when comparing amyloid with tangle, synuclein and TDP-43 load [57]. These findings have been replicated by other groups [58, 59], supporting the use of CSF neurogranin as a biomarker to detect synaptic dysfunction in response to A β pathology. Recent data from the BioFINDER study indicates that CSF neurogranin increases significantly shortly after A β accumulation, which further supports the idea that synaptic loss occurs early in AD patients [36]. Although initially thought to be AD-specific, a recent study demonstrated increased CSF neurogranin in CJD patients compared with cognitively unimpaired individuals and AD patients [60]. Other emerging biomarkers reflecting synaptic degeneration include synaptosomal-associated protein 25 (SNAP-25) [58, 61] and synaptotagmin-1 (SYT-1) [62], which are increased in AD and MCI compared with HC. Growth-associated protein 43 (GAP-43) is also increased in AD compared to controls, and reflects amyloid and tau pathology, and seems to be AD-specific [63], although transient increases can be found after stroke [64]. Another emerging biomarker is neuronal pentraxin 2 (NPTX-2), which in one pilot study showed lower levels in AD, as well as in individuals with cognitive impairment [65].

Blood

No studies measuring neurogranin in plasma have yet been able to differentiate AD from HC, due to relatively high peripheral expression confounding a potential difference [66]. Exosomal enrichment might overcome this issue. Studies examining neuronally derived exosomal neurogranin in plasma have been able to separate HCs from AD [67, 68], and stable MCI from MCI converting to AD [68], but more replication studies in this field are needed.

ALPHA-SYNUCLEIN PATHOLOGY

Misfolding of the α -synuclein protein plays a major role in the development of common NDDs, such as PD and DLB, and other neuropathologically related disorders. It is also the main constituent of Lewy bodies, the signature protein inclusion found in PD and DLB. Therefore, much of the research on diagnostic markers for these diseases has been focused on identifying pathology-specific forms of the protein.

CSF

It is relatively easy to measure total α -synuclein concentration in CSF but its level does not correlate with Lewy body pathology. Instead, it is argued that high levels are a marker of neurodegeneration [69]. Additionally, a recent study in ADAD mutation carriers as well as sporadic AD patients demonstrated increased levels of total α -synuclein and suggests that α -synuclein is linked to AD pathophysiology through the *APOE* ϵ 4 allele. Furthermore, total α -synuclein concentration in CSF increases if there is blood contamination of the sample, due to high expression in red blood cells, which further reduces its utility [70]. However, studies measuring oligomers and post translational modifications of α -synuclein have shown at least slightly positive results, but still not satisfactory enough for clinical use (for review, see [69]). Nonetheless, total α -synuclein in CSF might be used as a prognostic marker of motor progression in PD [71-73], and increased levels as a marker of synaptic degeneration [74]. Further, the idea of α -synuclein oligomers spreading in a prion-like manner has sparked the idea that seeding aggregation assays, such as real-time quaking induced conversion (RT-QuIC) or protein-misfolding cyclic amplification (PMCA) could be methods to qualitatively detect pathological forms of α -synuclein in CSF [75]. Studies analyzing CSF with RT-QuIC have been able to distinguish synucleinopathies from non-synucleinopathies with excellent diagnostic accuracy, detecting DLB and multiple system atrophy with 100% and 80% sensitivity, respectively [76]. Another study did a similar comparison and was able to discriminate between synucleinopathies (PD and DLB), and non-synucleinopathies with 100% specificity and 93 % sensitivity [77]. Unsurprisingly, these assays did not discriminate between different synucleinopathies [77]. However, panels combining different biomarkers may be a way to differentiate PD from atypical parkinsonian disorders, with one combination reaching an AUC of 0.95 [74]. In conclusion, results from α -synuclein aggregation assays are encouraging, but need to be validated in larger cohorts. There is also a need find reliable biomarker signatures to distinguish between different synucleinopathies.

Blood

Because of the high risk of contamination of plasma and serum samples with α -synuclein from RBC [70], it is not a surprise that total α -synuclein has yielded disappointing results as a diagnostic marker in whole blood, serum and plasma. However, better discriminatory power has been seen when instead measuring oligomeric or post-translationally modified α -synuclein (for review, see [69]). The encouraging results from aggregation assays in CSF may prove to be effective in other biofluids.

GLIAL ACTIVATION AND NEUROINFLAMMATION

Neuroinflammation, as well as activation of microglial cells and astrocytes, are key features of most NDDs, with most research having been performed in AD. During the last decade, it has been debated whether neuroinflammation and astrogliosis is an important driver of neurodegeneration, or a side effect of the accumulation of amyloid and tau.

CSF

A number of candidate markers have been examined in relation to inflammation/astroglial activation in NDDs, of which YKL-40, a glycoprotein expressed in both astrocytes and microglia, has proven to be maybe the most promising. Several cross-sectional, as well as longitudinal, studies in the past years have shown that CSF YKL-40 levels are modestly increased in patients with AD and FTD, but are relatively low in DLB, providing evidence for YKL-40 as a marker of neuroinflammation in the mentioned diseases [78, 79]. A study in the ADNI cohort showed conflicting results, with no significant separation of AD from HCs, but small sample sizes might not have given the power to detect a difference [58]. However, two recent longitudinal studies show that higher levels of YKL-40 in CSF correlate with cognitive decline as well as CSF tau levels [80, 81], and also with cortical thickness in CSF A β -positive individuals [81]. Increased levels in ADAD mutation carriers compared to non-carriers seem to appear 15 years before symptomatic disease is expected, together with markers of synaptic degeneration and neuronal injury [82]. A recent longitudinal study by Villar-Piqué *et al.* show higher plasma YKL-40 concentration among CJD patients compared with other NDs and HCs. This suggests that YKL-40 may be used as a marker of disease progression in CJD [83]. Lower levels of CSF YKL-40 compared to HCs have been observed in PD [84]. Other markers expressed in microglial cells that show promise is soluble triggering receptor expressed on microglial cells 2 (sTREM2), the secreted proteolytic degradation product of TREM2, which has been found at higher concentration in CSF from AD patients compared with HCs [85] with increases being observed already in individuals with reported subjective cognitive decline (SCD) [80]. Recently, the concentrations of CSF sTREM2 have been shown to be increased also in early symptomatic stages of sporadic AD [86, 87]. Interestingly, A β pathology and tau-related neurodegeneration may impact levels of CSF sTREM2 differently [87]. Data from the DIAN study of ADAD mutation carriers demonstrates that sTREM2 concentration in CSF increases in mutation carriers compared to non-carriers before symptomatic disease is expected, and after A β and T-tau [88]. Moreover, it has been shown that the concentrations of CSF sTREM2 vary between different disease-associated TREM2 genetic variant carriers [86, 87].

However, the highest CSF levels are found in patients with the autoimmune disease multiple sclerosis (MS) [89]. Other promising markers are monocyte chemoattractant protein (MCP-1) in CSF [80] and cytokine markers [81]. In conclusion, neuroinflammatory markers might be useful to pinpoint neuroinflammation in NDDs, which potentially could be of use as patient selection tools for clinical trials, in addition to core protein markers, or as a reflection of disease progression.

Blood

Some data suggest that YKL-40 is increased in plasma among patients with early AD [78, 90], but more studies are needed to confirm this finding. One study reported that higher plasma MCP-1 levels are associated with disease severity and faster cognitive decline in AD patients [91]. While sTREM2 is highly abundant in plasma and serum sTREM2 levels do not differ between multiple sclerosis, other inflammatory neurologic diseases and non-inflammatory controls [92]. Kleinberger et al. found no difference in plasma sTREM2 between healthy controls, AD and FTD [93]. Furthermore, a recent study indicates that there are no changes in plasma sTREM2 in *TREM2* rare variants carriers (Ashton NJ et al., 2019 *In Press*). Studies suggesting such a change in *TREM2* mRNA/protein levels among non-mutated *TREM2* AD patients should therefore be carefully interpreted [94].

TDP-43 PATHOLOGY

Cytoplasmic inclusions of TDP-43 is implicated in a majority of patients with the motor neuron disease amyotrophic lateral sclerosis (ALS), FTD [95] as well LATE, a neuropathological disease entity clinically indistinguishable from AD [6].

CSF and Blood

Over the last ten years, proteomic studies targeting TDP-43 pathology have only generated modest results. CSF TDP-43 can be measured but is subjected to contamination with peripheral expression, and correlation with neuropathological findings are weak [96]. Both phosphorylated [97, 98] and total [99] plasma TDP-43 can be measured and show weak or no correlation with neuropathological examination. Due to the difficulties separating peripheral from CNS-derived or pathological forms of TDP-43, more work needs to be done to develop a biomarker for TDP-43 pathology.

PRION PATHOLOGY

The conversion of the physiological, cellular form of prion protein (PrP^C) to the pathogenic, misfolded form of the protein (prion protein scrapie, PrP^{Sc}), together with extensive neuronal loss and spongiform appearance of the brain tissue at autopsy, characterizes this rare (incidence of 1 per million/year) group of fast progressing dementia, with CJD being the most common form.

CSF

Despite the difficulties of developing trustworthy specific immunoassays for the diagnosis of CJD and other prionopathies, the aggregation assay RT-QuIC has proven to be an effective method to detect PrP^{Sc}, which was first discovered in 2010 [100]. These results have since been replicated [101, 102], and according to a validation study, sporadic CJD was detected with a sensitivity of 92% and specificity of 100% using the method [103]. Among the patients who had a negative RT-QuIC result, 90% were positive using another diagnostic method [103].

Blood

The sensitivity of this assay triggers the thought of RT-QuIC as a potential blood test for prion disease. However, studies have shown that blood contamination decreases the performance of the CSF assay by effectively inhibiting the aggregation cascade [104], which has decreased the interest in developing a blood-based assay for the diagnosis of CJD.

NEURONAL INJURY

Neurofilament light (NFL) is the major biomarker for neuronal degeneration and is increased in neurodegenerative, inflammatory, traumatic and vascular conditions [105]. Another promising biomarker reflecting neuronal injury in NDDs is the neuronal calcium sensor protein visinin-like protein 1 (VLP-1/VILIP-1). Increased concentrations of CSF VLP-1 in AD patients compared with HCs are a well replicated finding [17]. Recent longitudinal studies report increased VLP-1 levels in amyloid-positive MCI and AD patients compared with amyloid-negative MCI and HCs [58, 106], and a longitudinal decrease in AD patients [58], possibly reflecting extensive neuronal damage. VLP-1 also predicts progression from MCI to AD [106]. An association with future cognitive decline, hippocampal atrophy among amyloid-positive MCI and AD was also found [106]. In contrast to NFL, VLP-1 also differentiates between AD and non-AD dementias [107, 108]. In conclusion, VLP-1 may be used in addition to NFL as a marker of neurodegeneration and disease progression in AD. Data on blood VLP-1 remains limited [107].

FUTURE DIRECTIONS

The rapid advancement in highly sensitive quantitative technologies has led to the promising translation of CSF biomarkers to blood and this includes A β , T-tau, p-tau and NFL. While studies on plasma A β IPMS platforms show great promise in terms of diagnostic accuracy, it is a method with complexity, especially considering the desired use of plasma A β as a screening tool with potential in primary care settings. Furthermore, no consensus cut-off values have been established yet, which is an important question to address. Additionally, there is still large inter-assay variability, and collaborative efforts are required to elucidate this. At AAIC 2019, there were different groups presenting plasma p-tau₁₈₁ assays, reflecting tangle pathology, which is much needed for enrollment in clinical trials and future primary care screening procedures. Standardized measurements for these two core proteins would facilitate cheaper and more reliable subject inclusion in much warranted clinical trials. However, as previously mentioned, further investigations into what tau measurements in biofluids actually tell us are warranted. Moreover, since the overwhelming proportion of academic studies is performed in white Caucasian populations, it is not known how the assays developed perform in individuals of other ethnic backgrounds. Since an increasing proportion of people with NDDs are non-Caucasians, it is important to investigate how the available assays perform in these populations. Another important matter that needs a solution is the absence of reliable diagnostic biomarkers for major NDDs, such as TDP-43 pathology in FTD and ALS, and α -synuclein pathology in PD and DLB. In conclusion, cooperative efforts are needed to validate the promising AD plasma assays available, as well as intensifying the search for biomarkers as reliable diagnostic tools for other NDDs.

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HIGHLIGHTS

- Protein biomarkers in cerebrospinal fluid now define Alzheimer's disease (AD).
- New ultrasensitive techniques have enabled promising replication in blood.
- This would enable the use of biomarkers to screen for AD in a primary care setting.
- The coexistence of pathologies complicates clinical trials and diagnostics of neurodegenerative diseases (NDDs).
- Biomarkers for NDDs that reflect other pathologies are greatly needed.

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This study analyzing CSF with RT-QuIC provide evidence that synucleinopathies can be distinguished from non-synucleinopathies with excellent diagnostic accuracy.

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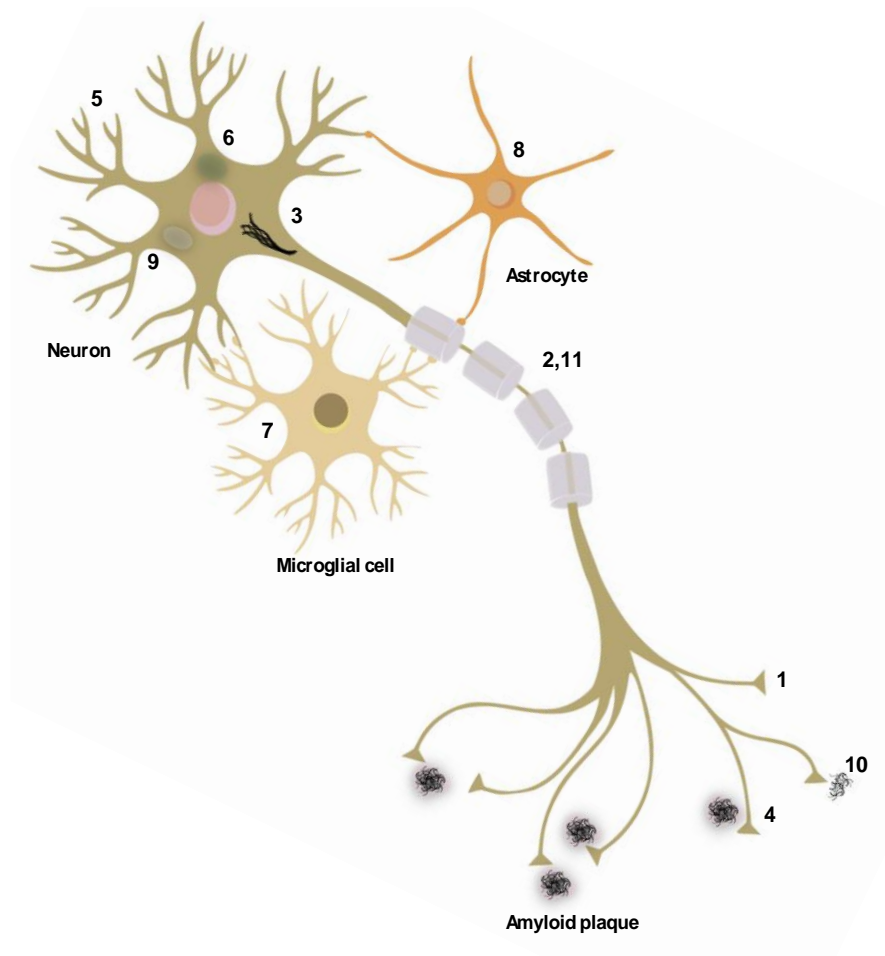
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Amyloid pathology

1. $A\beta_{42}$, $A\beta_{40}$, $A\beta_{38}$, $A\beta$ oligomers

Tau pathology

2. T-tau
3. P181-tau in neurofibrillary tangles

Synaptic degeneration

4. SNAP-25, SYT-1,
5. GAP-43, neurogranin, NPTX-2

α -Synuclein pathology

6. α -Synuclein inclusion

Glial activation and neuroinflammation

7. TREM-2, MCP-1, YKL-40, cytokines
8. YKL-40, cytokines

TDP-43 pathology

9. TDP-43 inclusion (no marker available)

Prion pathology

10. PrPSc aggregation

Neuronal injury

11. VLP-1

ers and localization of

Method	Biomarkers measured in plasma	Change in Amyloid (A) + VS A- group	Receiver Operating Characteristic (ROC) (A+ VS A- as diagnosed by PET and CSF)	Correlation of plasma biomarkers with PET (Mean Cortical Standard Uptake Value Ratio (mSUVR)) and CSF	Reference
IP-MS (LC/MS)	<ul style="list-style-type: none"> Aβ38 Aβ40 Aβ42 	<ul style="list-style-type: none"> Aβ42: ↓ in A+ group Aβ42/ Aβ40 ratio: ↓ in A+ group 	<p>PET Aβ42/ Aβ40 ratio: AUC= 0.88</p> <p>CSF (p-tau181/Aβ42) Aβ42/ Aβ40 ratio: AUC= 0.85</p>	<p>PET Aβ40/ Aβ42 ratio: r= -0,55</p> <p>CSF Plasma Aβ42/ Aβ40 ratio and CSF Aβ42/ Aβ40: r= 0,66</p>	[19]
IP-MS (MALDI-TOF)	<ul style="list-style-type: none"> Aβ40 Aβ42 APP669-711 	<ul style="list-style-type: none"> Aβ42: ↓ in A+ group Aβ40/ Aβ42 ratio: ↑ in A+ group APP669-711/ Aβ42 ratio: ↑ in A+ group Composite biomarker: ↑ in A+ group <p>Composite biomarker equals average of the normalized APP669/711: Aβ42 ratio and Aβ40/ Aβ42 ratio</p>	<p>PET Aβ42: AUC= 0.789 Aβ40/ Aβ42 ratio: AUC= 0.886 APP669-711/Aβ42 ratio: AUC= 0.861 Composite biomarker: AUC= 0.914</p> <p>CSF AIBL sub-group (n=46) Composite biomarker: AUC= 0.876</p>	<p>PET Aβ42: r= -0.484 Aβ40/ Aβ42 ratio: r= 0.626 APP669-711/Aβ42 ratio: r= 0.606 Composite biomarker: r= 0.678</p> <p>CSF Aβ42 and CSF Aβ42: r=0.408 Aβ40/ Aβ42 ratio and CSF Aβ42: r=-0.534 APP669/711:Ab42 ratio and CSF Aβ42: r=0.601 Composite biomarker and CSF Aβ42: r=-0.660</p>	[18]
ELISA	<ul style="list-style-type: none"> Aβ40 (total and free) Aβ42 (total and free) 	<ul style="list-style-type: none"> TAβ42/ TAβ40 ratio: ↓ in A+ group 	<p>PET TAβ40/ TAβ42 ratio AUC= 0.775</p>	<p>PET (mean global cortical amyloid SUVR) TAβ40/ TAβ42: r= -0,514</p>	[22]
ELECSYS	<ul style="list-style-type: none"> Aβ40 Aβ42 	<ul style="list-style-type: none"> Aβ42: ↓ in A+ group Aβ42/Aβ40 ratio: ↓ in A+ group 	<p>CSF Aβ42: AUC = 0.71 Aβ42/ Aβ40 ratio: AUC = 0.77 Aβ42, Aβ40: AUC = 0.80</p>	<p>CSF Plasma Aβ42 and CSF Aβ42: r=0.373 Plasma Aβ42/ Aβ40 ratio and CSF Aβ42/ Aβ40: r=-0.476 Plasma Aβ40 ratio and CSF Aβ40: r=0.100</p>	[25]
SIMOA	<ul style="list-style-type: none"> Aβ40 Aβ42 	<ul style="list-style-type: none"> Aβ42: ↓ in A+ group Aβ42/ Aβ40 ratio: ↓ in A+ group 	<p>PET Aβ42: AUC = 0.604 Aβ42/ Aβ40 ratio: AUC = 0.621</p> <p>CSF Aβ42: AUC = 0.655 Aβ42/ Aβ40 ratio: AUC = 0.683</p>	<p>PET Aβ42: r=-0.162 Aβ40: r=-0.012 Aβ42/ Aβ40 ratio: r=-0.167</p> <p>CSF Plasma Aβ42 and CSF Aβ42: r=0.274 Plasma Aβ42/ Aβ40 ratio and CSF Aβ42/ Aβ40: r=0.215 Plasma Aβ40 ratio and CSF Aβ40 : r=0.136</p>	[20]
SIMOA	<ul style="list-style-type: none"> Aβ40 Aβ42 	<ul style="list-style-type: none"> Aβ42: ↓ in A+ group Aβ42/ Aβ40 ratio: ↓ A+ group 	<p>PET Aβ42: AUC = 0.66 Aβ42/ Aβ40 ratio: AUC = 0.68</p> <p>CSF Aβ42: AUC = 0.66 Aβ42/ Aβ40 ratio: AUC = 0.77</p>	<p>CSF Plasma Aβ42 and CSF Aβ42: r=0.18 Plasma Aβ42/ Aβ40 ratio and CSF Aβ42: r=0.38</p>	[21]
IMR	<ul style="list-style-type: none"> Aβ42 	<ul style="list-style-type: none"> Aβ42: ↑ in A+ group 		<p>CSF Plasma Aβ42 and CSF Aβ42 r= -0.352</p>	[23]
IMR	<ul style="list-style-type: none"> Aβ40 Aβ42 	<ul style="list-style-type: none"> Aβ40: ↓ in AD (clinical diagnosis) Aβ42: ↑ in AD (clinical diagnosis) 	<p>PET Aβ42: AUC = 0.776</p>	<p>PET Aβ40: r=-0.585 Aβ42: r=-0.281</p>	[24]
	<ul style="list-style-type: none"> Exosome-bound Aβ42 	<ul style="list-style-type: none"> Exosome-bound Aβ42: ↑ in A+ group 		<p>PET Exosome-bound Aβ42: r=0.9002</p>	[29]
	<ul style="list-style-type: none"> Aβ – heterogeneous (monomers + oligomers/aggregates) and monomerized states 	<ul style="list-style-type: none"> ↑ self-standard ratio in AD compared with CU 		<p>PET Self –standard ratio (AMC): r=0.5511 Self –standard ratio (KIRAMS): r=0.4141</p>	[28]

Table 1. A summary of the results from studies investigating plasma amyloid as a biomarker of cerebral amyloidosis.