

Expert Review of Neurotherapeutics

Cerebrospinal fluid in the differential diagnosis of Alzheimer's disease: an update of the literature

Abstract

Introduction: The importance of cerebrospinal fluid (CSF) biomarkers in Alzheimer's disease (AD) diagnosis is rapidly increasing, and there is a growing interest in the use of CSF biomarkers in monitoring the response to therapy, especially in the light of newly available approaches to the therapy of neurodegenerative diseases.

Areas covered: In this review we discuss the most relevant measures of neurodegeneration that are being used to distinguish patients with AD from healthy controls and individuals with mild cognitive impairment, in order to provide an overview of the latest information available in the scientific literature. We focus on markers related to amyloid processing, markers associated with neurofibrillary tangles, neuroinflammation, neuroaxonal injury and degeneration, synaptic loss and dysfunction, and markers of α -synuclein pathology.

Expert opinion: In addition to neuropsychological evaluation, core CSF biomarkers ($A\beta_{42}$, t-tau, and p-tau181) have been recommended for improvement of timely, accurate and differential diagnosis of AD, as well as to assess the risk and rate of disease progression. In addition to the core CSF biomarkers, various other markers related to synaptic dysfunction, neuroinflammation, and glial activation (neurogranin, SNAP-25, **Nfl**, YKL-40, TREM2) are now investigated and have yet to be validated for future potential clinical use in AD diagnosis.

Keywords: Alzheimer's disease; Amyloid-beta; Biomarkers; Cerebrospinal fluid; Diagnosis; Neuroaxonal injury; Neuroinflammation; α -Synuclein; Synaptic dysfunction; Tau

Article highlights

- Besides decreased levels of $A\beta_{42}$ in CSF, CSF $A\beta_{40}$, $A\beta_{37}$ and $A\beta_{38}$ may help in distinguishing AD from healthy controls and other dementias, such as FTD and LBD.
- CSF $A\beta_{42}/A\beta_{38}$ and $A\beta_{42}/A\beta_{40}$ ratios exhibit enhanced performance in distinguishing AD from other forms of dementia, including PD, LBD, subcortical VaD and FTD.
- Elevated CSF levels of p-tau (p-tau181 and p-tau231) are a valuable marker in differentiating AD from other types of dementia.
- The p-tau217 exhibits higher accuracy than p-tau181 and p-tau231 in distinguishing AD dementia from non-AD.
- CSF p-tau/ $A\beta_{42}$ ratio could be accurate predictor of conversion from MCI to AD dementia.
- NfL provides important information about the progress of neurodegeneration and should be used as a biomarker in AD, but not as a biomarker of AD.
- CSF VILIP-1 levels are significantly increased in AD and MCI patients compared to controls, and correlates well with the progression and pathology of AD.
- Elevated CSF Ng and SNAP-25 levels are found in patients diagnosed with AD, even in prodromal phase of the disease.
- Individuals with AD exhibit higher levels of glycoprotein YKL-40 in CSF and plasma compared to healthy controls.
- AD pathology is associated with elevated CSF sTREM2 levels, especially in the case of tau-related neurodegeneration.
- Increased CSF α -syn levels are present in patients with MCI and AD, and they correlate with disease progression and/or severity of cognitive decline.

1 Introduction

Alzheimer's disease (AD) is a chronic progressive neurodegenerative disease with an irreversible but slow time course, and it is the most frequent of all types of dementia (around 60-70% cases). It is characterized with cognitive deterioration, various neuropsychiatric symptoms, and behavioral problems, and in the later stage with inability to perform daily living activities [1]. The main neuropathological features in AD are the accumulation of extracellular amyloid β ($A\beta$) plaques and the intracellular accumulation of hyperphosphorylated tau proteins as neurofibrillary tangles, that result in progressive neurodegeneration and cerebral atrophy. The detailed description about the AD pathophysiology was reviewed recently [2]. Pathophysiology of AD is described with different hypotheses: cholinergic hypothesis, amyloid cascade hypothesis and hyperphosphorylation of tau proteins in the brain [3]. All hypotheses are confirmed with some pathological or pharmacological findings: a decreased acetylcholine concentration in the brain is associated with neuronal loss and is responsible for the memory loss, cognitive deterioration and AD development, and this hypothesis is confirmed by the clinical efficacy of the acetylcholinesterase inhibitors. Amyloid cascade hypothesis is the most accepted, suggesting that the primary cause of AD is the extent accumulation of $A\beta$ peptides and $A\beta$ plaque deposition in the brain. After the enormous and long-term efforts to develop a treatment, some new drugs targeting $A\beta$ have shown satisfactory results [4,5]. A common hypothesis is also the hyperphosphorylation of tau proteins, due to the excessive accumulation of the amyloid proteins in the brain tissue. The hypothesis is based on abnormal changes in tau proteins which result in tau dysfunction, and these hyperphosphorylated tau proteins shape aggregates (i.e. neurofibrillary tangles) that are deposited within the neurons, with consequent induction of neuronal damage and negative effects on neuronal function [3]. The full description of the AD pathophysiology regarding the $A\beta$ plaques and neurofibrillary tangles is provided in the text in paragraphs 2.1. (markers of beta-amyloid accumulation and amyloid processing) and 2.2. (markers of tau pathology), while other pathophysiological features of AD are described further in the text in the paragraphs 2.3-2.6 (markers of neuroaxonal injury and degeneration, markers of synaptic dysfunction, markers of neuroinflammation, markers of α -synuclein pathology).

Currently there is no cure for AD, no disease-modifying treatment, and no therapeutic strategy exists to prevent AD or reverse disease progression, while few classes of medication (i.e. acetylcholinesterase inhibitors and non-competitive NMDA antagonist) can only slow AD progression, with modest benefit on cognition [1]. Medication used to treat symptoms in AD include donepezil (reversible non-competitive acetylcholinesterase inhibitor), galantamine (reversible, competitive acetylcholinesterase inhibitor and modulator of nicotinic acetylcholine receptor), rivastigmine (acetylcholinesterase and butyrylcholinesterase inhibitor) and memantine (non-competitive NMDA antagonist) [1]. However, recently two additional treatment strategies have been approved by the U.S. Food and Drug Administration (FDA) for the treatment of AD, aducanumab [4] and lecanemab [5]. Both drugs are based on a humanized IgG1 monoclonal antibody that targets aggregated $A\beta$ and both approaches have been shown to reduce $A\beta$ burden measured with PET [6]. Lecanemab exhibits stronger binding to $A\beta$ protofibrils, while aducanumab has greater affinity for highly aggregated $A\beta$ fibrils [6]. In contrast to aducanumab, the efficiency of lecanemab was found to be more consistent across different studies [7–9].

The National Institute on Aging–Alzheimer's Association (NIA-AA) in 2018 shifted its guidelines of the definition of AD from the syndromal entity to biologically based entity and described AD as “underlying pathologic processes that can be documented by postmortem

examination or in vivo by biomarkers" [10]. This definition should be used for the research framework, and not for clinical practice [11]. The "A/T/N" classification scheme was proposed in order to categorize AD biomarker findings into a format which is easy to understand and use [11]. The classification included 7 major AD biomarkers subdivided into 3 categories: "A" ($A\beta$ plaques i.e. biomarkers detected with cortical amyloid PET ligand binding or decreased cerebrospinal fluid (CSF) $A\beta_{42}$), "T" (related to aggregated tau (neurofibrillary tangles), i.e. increased CSF phosphorylated tau (P-tau) and cortical tau PET ligand binding), and "N" (biomarkers of neurodegeneration or neuronal injury) [11]. The "N" biomarkers include CSF T-tau, FDG PET hypometabolism, and brain atrophy detected with MRI, but are open to other (novel) markers of neurodegeneration. This classification points out the importance of separating biomarkers related to neurofibrillary tangles from markers associated with neuronal injury/neurodegeneration since this approach might help differentiate neuronal AD from non-AD causes [11]. This shift in definition was suggested since these neuropathological changes (β -amyloid plaques and neurofibrillary tau deposits, evaluated as biomarkers) define AD, and might offer more precise characterization and improve the knowledge of the order of neurobiological events that cause AD, however, in addition, AD should be assessed also in different stages across its entire spectrum [11]. Therefore, the difference is that AD is not only defined by its clinical symptoms (cognitive deterioration that affects thinking, remembering, and reasoning, and behavioral changes that interfere with daily life and activities), but also with characteristic neuropathologic changes that can be assessed with biomarkers in vivo and postmortem examination. However, when evaluating biomarkers, besides sex-specific contributions to AD risk biomarkers [12], racial differences should also be considered. The African American individuals were suggested to have lower levels of CSF t-tau and p-tau181 compared to white individuals, suggesting that in these individuals ATN biomarkers must be controlled for race [13].

Regarding the clinical perspective, AD diagnosis is met when all criteria defined by The American Psychiatric Association's Diagnostic and Statistical Manual (DSM-5) [14] for dementia are present, and the neuropsychiatric symptoms (cognitive decline associated with reduced ability for reasoning or thinking, problems with memory and in social skills, behavioral abilities) occur [15]. Problems in cognition are associated with two or more of the following domains: acquisition and recall of information, reasoning and judgment in complex tasks, visuospatial ability, language function, and normal personality or behavior [15]. AD can be divided in different clinical stages [16]. The first is a pre-clinical or pre-symptomatic stage, lasting for several years or more, with characteristic mild memory loss and early pathological changes in cortex and hippocampus, but without any functional impairment in the daily activities and there are no clinical symptoms of AD [16]. After that a mild or early stage of AD occurs, with development of particular symptoms (i.e. changes in mood, problems in concentration and memory, disorientation of place and time, and sometimes depression) [16]. A moderate stage of AD is characterized with elevated loss of memory, troubles in recognizing family and friends, deficit in impulse control, and problems in reading, writing, and speaking, while a severe stage of AD or a late-stage of AD is associated with advanced functional and cognitive deterioration, where patients are not able to recognize their family, have problems in swallowing and urination, and this is an end-stage of AD where affected patients die from various complications [16]. Since AD is presented as a continuum, different stages associated with time course of AD exist, depending on the working group that defines them (International Working Group or NIA-AA), and can be divided into: asymptomatic at risk or preclinical AD; prodromal phase or AD with MCI; mild AD dementia or AD with mild dementia; moderate AD dementia or AD with moderate dementia; and the last severe AD dementia or AD with severe

dementia [17]. All these stages are characterized with specific biomarkers, and therefore evaluation of these biomarkers in early stages in asymptomatic patients or those with MCI might offer personalized approach and therapy that targets specific pathological processes and corresponding biomarkers [2].

In the possible prevention, treatment and reduction of the symptoms of AD, early screening and accurate diagnosis are the most important. Biomarkers should be used for the early diagnosis of AD, and this approach offers an improved therapeutic strategy to slow the progression of disease and treat or reduce the symptoms [18]. Following recent recommendations of the International Working Group, biomarkers should detect individuals in early stages who are at risk for progression to AD dementia, or to AD, and will differentiate various types of dementia and different AD phenotypes and assist in identifying those at risk for symptomatic AD [18,19]. However, the prediction or diagnosis of AD should not be based exclusively on AD biomarkers, but should be combined with the clinical assessment [18]. The diagnosis of AD already starts in primary health care with taking patients' medical history, which is afterwards complemented by physical examination, laboratory tests and cognitive screening (Figure 1). The importance of including both neuroimaging and CSF biomarkers is in ruling out AD as the underlying cause of cognitive dysfunction (Figure 1), in enabling earlier, more accurate and differential diagnosis of AD, and in enabling personalized management and treatment of AD [20].

2 CSF biomarkers in the diagnosis of AD

The importance of CSF biomarkers in the diagnosis of AD is rapidly increasing, and this interest is also growing in terms of monitoring the response to therapy, especially in the light of newly available approaches to the therapy of neurodegenerative diseases. Determination of CSF amyloid beta ($A\beta$), total tau (t-tau), and phosphorylated tau (p-tau) levels has been included in both clinical and research diagnostic criteria [18,19,21].

The existing literature is mostly focused on biomarkers that are associated with AD pathogenic process, and the main clinical value of these biomarkers is to facilitate the diagnosis of AD and help the clinician to discriminate between AD associated phenotypes and non-AD pathologies. Diagnostic biomarkers in AD focus on three main pathological features of AD which reflect the process underlying AD: accumulation of $A\beta$, hyperphosphorylation of tau, and neuronal degeneration (less specific markers of AD) [11]. CSF biomarkers that have been researched the most and have shown the best biomarker properties so far include $A\beta_{42}$, $A\beta_{40}$, p-tau₁₈₁, p-tau₂₁₇, and t-tau. These markers accurately identify the pathological changes associated with AD, even in early stages of the disease (asymptomatic and prodromal stages), and they have potential to predict cognitive decline [22]. Markers associated with $A\beta$ accumulation and amyloid processing and the markers of tau pathology are useful for the differentiating AD patients from those with non-AD dementia which will be discussed in more details in the following sections of this review. Some clinical practices have replaced t-tau as a measure of neurodegeneration with neurofilament light chain (NfL) because of its high sensitivity [23]. Other CSF biomarkers which are currently being researched, and will be discussed in the further text, have a better potential to one day be used as biomarkers for staging of disease and prognosis, less for the differential diagnoses [18].

This review concentrates on the most relevant measures of neurodegeneration that are being determined in order to distinguish patients with AD from healthy control subjects and individuals with mild cognitive impairment (MCI) in order to provide an overview of the latest information available in the scientific literature. The review focuses on markers related to

amyloid processing, markers of tau pathology, neuroinflammation, neuroaxonal injury and degeneration, synaptic loss and dysfunction, and markers of α -synuclein pathology (Figure 2).

2.1 Markers of beta-amyloid accumulation and amyloid processing

A significant progress in AD research was the identification of $A\beta$ as the primary protein component of amyloid plaques, formed through the enzymatic breakdown of its precursor, the amyloid precursor protein (APP). In the amyloidogenic pathway, APP is cleaved by β -secretase, resulting in the production of sAPP β and β CTF 99 fragments [24]. Subsequently, β CTF 99 is cleaved by γ -secretase, leading to the generation of $A\beta$ peptides of various lengths, predominantly $A\beta_{42}$ and $A\beta_{40}$, with the former being much more prone to aggregation [25] and the formation of amyloid plaques associated with AD. In contrast, the nonamyloidogenic pathway involves cleavage by α -secretase and production of sAPP α and α CTF 83 fragments, while further processing by γ -secretase generates peptides like p3, which is less implicated in AD pathology [26].

The CSF levels of $A\beta_{42}$, when combined with total tau (t-tau) and phosphorylated tau (p-tau), form the recognized CSF pattern used to diagnose AD, with $A\beta_{42}$ demonstrating the highest diagnostic accuracy [27]. The sensitivity for predicting progression to AD in patients with MCI was found to be 95%, and the specificity 87% [28]. For distinguishing AD from other conditions, the sensitivity of CSF $A\beta_{42}$ was 85% and the specificity 42% in case of Lewy body disease (LBD), 85% and 77% in case of frontotemporal dementia (FTD) [27], and 77% and 80% for vascular dementia (VaD) [29]. Moreover, evidence suggests that $A\beta_{42}$ levels may serve as a predictor of disease progression in individuals with normal cognition and those with MCI [27]. Levels of CSF $A\beta_{42}$ were significantly reduced in AD patients compared to normal individuals [30]. This reduction is attributed to increased deposition of amyloid plaques in the brain, leading to decreased levels of $A\beta_{42}$ in CSF. In previous studies, reduced CSF levels of $A\beta_{42}$ have also been observed in other non-AD disorders, such as FTD, LBD, and VaD [30]. Nevertheless, these observations could be linked to mixed pathology, suggesting the existence of various pathological conditions or overlapping characteristics among the aforementioned diseases. According to the recent study, it is proposed that the p-tau/ $A\beta_{42}$ ratio could serve as a valuable tool in distinguishing AD from FTD with primary language difficulties [31]. However, the same ratio does not effectively differentiate AD from behavioral variant or FTD nor from FTD as a collective entity [31]. In addition, CSF $A\beta_{42}$ could be a differentiating marker for the detection of prodromal AD in clinically diagnosed amnesic MCI patients [32]. Conversely, another study found that, compared to AD, cerebral amyloid angiopathy (CAA) showed similar levels of $A\beta_{42}$, but lower levels of $A\beta_{40}$ [33].

Previous study has reported varied findings regarding CSF $A\beta_{40}$ levels [34]. The reports showed either decreased, unchanged, or elevated CSF $A\beta_{40}$ levels in AD compared to other forms of dementia [34]. Recent research highlighted a significant age-independent rise in CSF $A\beta_{40}$ levels in AD, along with a positive association between CSF $A\beta_{40}$ and p-tau₁₈₁ concentration, even in non-AD individuals, suggesting that initial amyloid peptide levels may serve as a risk factor for sporadic AD [34]. On the other hand, diminished CSF $A\beta_{40}$ levels could indicate alternative conditions such as FTD, CAA [34], normal pressure hydrocephalus [35] and multiple sclerosis [36].

Besides $A\beta_{42}$ and $A\beta_{40}$, $A\beta_{37}$ and $A\beta_{38}$ may help in distinguishing AD from other dementias, such as FTD and LBD [30]. The noticeable decrease in CSF $A\beta_{42}$ levels was reported in AD, while in LBD the CSF levels of $A\beta_{38}$, $A\beta_{40}$ and $A\beta_{42}$ were found to be decreased [37]. This indicates that amyloid metabolism is altered in LBD, even when there is no concurrent

presence of AD pathology. Furthermore, differentiating AD from CAA using CSF biomarkers is challenging due to symptom overlap between these diseases [38]. It was suggested that adding $A\beta_{38}$ and $A\beta_{43}$ to standard AD biomarkers could improve this differentiation [39], while others found no enhancement in distinguishing between AD and CAA with their inclusion [40]. The significant improvement in AD diagnosis was achieved through calculating the CSF $A\beta_{42}/A\beta_{40}$ ratio [41]. The CSF $A\beta_{40}$ initially showed limited potential as a stand-alone biomarker. However, it quickly became apparent that normalizing $A\beta_{42}$ levels to the total $A\beta$ quantity, represented by the most abundant isoform $A\beta_{40}$, yields superior diagnostic performance compared to $A\beta_{42}$ alone [42]. Consistent $A\beta_{40}$ levels were observed across AD patients, non-AD patients and controls, but the reduction in $A\beta_{42}$ levels significantly elevated the $A\beta_{42}/A\beta_{40}$ ratio [27]. Subsequent study revealed notable differences in $A\beta_{42}/A\beta_{38}$ and $A\beta_{42}/A\beta_{40}$ ratios in AD, suggesting superior diagnostic potential compared to individual biomarkers [27]. Further, CSF $A\beta_{42}/A\beta_{38}$ and $A\beta_{42}/A\beta_{40}$ ratios exhibited enhanced performance in distinguishing AD from other forms of dementia, including Parkinson's disease dementia (PD), LBD and subcortical VaD [43]. In addition, it was proposed that the $A\beta_{42}/A\beta_{40}$ ratio is highly valuable for differentiating between AD and LBD patients, especially during the prodromal stage when clinical diagnosis proves to be particularly challenging [44]. Moreover, recent study showed that the $A\beta_{42}/A\beta_{40}$ ratio outperformed $A\beta_{42}$ alone in distinguishing AD from FTD [45].

In addition to these findings, CSF sAPP α and sAPP β have been proposed as potential novel CSF biomarkers for AD and several other neurodegenerative conditions [46]. Their effectiveness has not met expectations, often yielding conflicting results, with limited studies examining their utility in distinguishing between different neurodegenerative diseases beyond comparisons with healthy elderly controls [47,48]. Significant elevations in both sAPP α and sAPP β levels in individuals in the MCI-AD group compared to those in the MCI-others group were reported [47]. The same group also verified a strong correlation between levels of sAPP α , sAPP β , p-tau, as well as t-tau, suggesting potential pathological connections between tau and sAPPs [48]. This correlation is significant as the increase in p-tau and t-tau levels is believed to indicate the neurodegenerative alterations linked to AD [48]. Finally, the study by Alcolea et al. offers pathological confirmation that low levels of sAPP β and high levels of chitinase-3-like protein 1 (CHI3L1 or YKL-40) in the CSF are linked to FTD [49]. Thus, these biomarkers may be valuable, especially in specific clinical situations where FTD is suspected.

2.2 Markers of tau pathology

P-tau protein levels along with $A\beta_{42}$ levels are known as "core CSF biomarkers" and are assumed to have the highest diagnostic accuracy for the early diagnosis of AD [50–52]. The NFTs are considered as second neuropathological hallmarks of AD, after A β plaques [53]. NFTs are composed of a highly-phosphorylated form of the microtubule-associated protein tau [54]. Tau protein is mainly present in axons and plays an important role in connecting microtubules and controlling axonal length and stability [55]. Abnormal phosphorylation of tau proteins causes detachment of tau from microtubules, degradation of microtubules, which affects axons, and ultimately leads to neuronal death [56]. Injury and degradation of axons and neuronal cell death, lead to the release of tau protein to CSF, which is reflected as t-tau levels [57,58]. Consequently, t-tau levels indicate the degree of neuronal loss and neurodegeneration in AD [59].

A key component of NFTs in AD pathology, and more accurate biomarker for differentiating AD from other dementias in contrast to t-tau, is p-tau. Recent study revealed at least 59 different p-tau phosphorylation sites which could be related to AD using mass spectrometry [60]. CSF p-tau levels have demonstrated strong prognostic accuracy in AD, especially in predicting cognitive decline in patients with AD and MCI [61,62]. Furthermore, the elevated levels of p-tau in AD, compared to other neurodegenerative conditions, such as VaD, FTD, progressive supranuclear palsy (PSP) or corticobasal syndrome (CBS), make it a valuable marker in differentiating AD from other types of dementia [63,64]. Several p-tau species have been reported to be increased in the very early stages of AD such as p-tau₁₈₁, p-tau₂₃₁ and p-tau₂₁₇ [65]. P-tau₁₈₁ is one of the most studied variants of phosphorylated tau protein, and it is considered a gold standard for AD diagnosis [66]. It is reported to be elevated in patients with MCI and AD continuum [67,68]. Even though p-tau₁₈₁ is elevated in AD, in other types of dementia, such as FTD, it is significantly decreased compared to healthy controls [69,70]. However, p-tau₁₈₁ is not significantly different between progressive supranuclear palsy, corticobasal degeneration, or other variants of FTD [70]. Another variant of p-tau is p-tau₂₃₁ that has been reported to be elevated in patients with AD and MCI, compared to controls [71,72]. Previous studies also demonstrated the role of p-tau₂₃₁ in differentiating AD patients from FTD, VaD and LBD with the sensitivity of 90.2% and with 80% of specificity [73]. More recent research has shown that p-tau₂₁₇ exhibited higher accuracy than p-tau₁₈₁ and p-tau₂₃₁ in distinguishing AD from non-AD dementia [74,75]. In a study conducted by Barthélemy and colleagues, p-tau₂₁₇ was a better predictor of A β positivity than p-tau₁₈₁ [74]. Compared to CSF p-tau₁₈₁, CSF p-tau₂₁₇ showed a stronger correlation with CSF A β ₄₂ and with A β and tau-PET [76]. Similarly, another study also reported better correlation of p-tau₂₁₇ in differentiating AD from other neurodegenerative diseases with a 91% sensitivity and specificity [77]. Additionally, in comparison to p-tau₁₈₁, p-tau₂₁₇ had a 90% accuracy rate in separating AD from tauopathies such as Pick disease, progressive supranuclear palsy, and corticobasal degeneration [78]. These studies suggest that p-tau₂₁₇ is the most accurate tau biomarker for AD in both pre-clinical and advanced stages.

Although CSF t-tau and p-tau are well recognized biomarkers for differentiating AD from other dementias, their diagnostic value is significantly enhanced when measured in combination with A β ₄₂ [22,79–81]. Furthermore, the combination of tau and A β markers was demonstrated to be useful in predicting disease progression [82,83]. A recent study revealed that CSF p-tau/A β ₄₂ ratio could be accurate predictor of conversion from MCI to AD dementia, with 82.9% sensitivity and 90% specificity [84]. It was previously reported that individuals with FTD exhibited the highest levels of A β ₄₂ and the lowest levels of t-tau and p-tau in FTD, whereas AD patients showed the highest levels of t-tau and p-tau, and the lowest levels of A β ₄₂ and A β ₄₂/p-tau ratios [85]. More recent study aimed to determine the CSF levels of tau and A β for distinguishing FTD from AD [31]. This study reported that the p-tau/A β ₄₂ ratio might be beneficial in distinguishing between AD and FTD characterized by primary language impairments, but was not effective in discriminating AD from the behavioral variant of FTD, or from FTD as a collective group [31]. Therefore, investigating and gaining a deeper understanding of the role of the tau protein in the mechanisms and underlying pathology of AD could enhance diagnostic and prognostic accuracy, particularly in distinguishing AD from other forms of dementia.

2.3 Markers of neuroaxonal injury and degeneration

Recently, the question has been raised whether AD can be considered as an axonal degeneration disease [86]. Both key factors in the pathogenesis of AD, tau and A β , have a predominant expression and an important role in the physiological functions of axons. When microtubules are broken or other axonal injuries occur, A β and tau may be abnormally modified and the result is deteriorated neuroaxonal damage. Neuroaxonal injury and consequential synaptic dysfunction are key features of AD, and synaptic loss is closely associated with cognitive decline in the disease [87]. One of the earliest changes that occur in early AD is axonal dystrophy which is associated with extracellular depositions of A β and has been observed to contribute to synaptic alterations occurring in AD [88]. As previously mentioned, t-tau levels represent a general signal of neurodegeneration [65]. Different studies reported a higher concentration of t-tau levels in CSF of AD patients in comparison to healthy subjects [89–91]. Since CSF levels of t-tau protein are believed to reflect the extent of neuronal damage, it is hypothesized that very high CSF t-tau levels, compared to moderately elevated levels, may correspond to differences in the degree of cortical atrophy and various clinical subtypes of AD [91]. Previous studies reported a faster rate of clinical progression in AD patients with high CSF t-tau levels [92,93]. Recent studies also reported elevated CSF t-tau levels in rapidly progressive AD (rpAD) [94] and in patients with atypical AD clinical phenotypes [91]. High concentration of CSF t-tau levels may not be very specific for AD, since high levels were observed in other types of dementia such as VaD [95] and FTD [95,96]. Furthermore, fluctuations in CSF t-tau levels also occur in cerebral ischemia [97], hemorrhage and seizures [98], as well as in cases of encephalitis and acute neurodegenerative diseases such as Creutzfeldt-Jakob disease (CJD) [99]. As a consequence of severe neuronal damage in CJD, studies reported t-tau levels are much higher in CJD than in AD.

Neurofilament light chain (Nfl) is one of four subunits of neurofilaments, which are proteins that are located in the neuronal cytoplasm that help maintain the structural stability of neurons and enable the growth of axons [100]. Due to its presence in neurons, when neuroaxonal damage occurs, Nfl levels increase in interstitial fluid and consequently, in CSF and plasma [101]. Therefore, the Nfl level is increased in a whole variety of neurological disorders, including neurodegenerative, inflammatory, traumatic and cerebrovascular diseases and is not specific to the neurodegenerative changes present in AD [102]. Although nonspecific, Nfl provides important information about the progress of neurodegeneration and should be used as a biomarker in AD, but not as a biomarker of AD [66]. The presence of Nfl in CSF and plasma correlates strongly, but concentrations in CSF are much higher and that is why it is the sample of choice for clinical use [103]. The disadvantage of CSF sampling is the invasiveness of the method and, consequently, the difficult implementation of longitudinal studies. With the development of ultra-sensitive techniques for the determination of biomarkers in plasma, such as Single molecule array (SIMOA), the determination of Nfl from plasma has become much more accurate. Measuring Nfl from plasma would solve the problem of invasiveness of sampling and enable longitudinal studies [104]. There is an increasing number of studies that indicate that both CSF and plasma Nfl may serve as diagnostic, prognostic and monitoring biomarkers in differential diagnosis between neurodegenerative diseases, including AD, and nondegenerative disorder, highlighting the Nfl as one of the most promising biomarkers to be used in clinical and research settings in the future [100]. This conclusion is also supported by the results of a recent study that reported a good accuracy for plasma Nfl levels in distinguishing between ATN+ and ATN– subjects in the group of patients with subjective cognitive decline (AUC=0.815) and MCI patients (AUC=0.818) [105].

Visinin-like protein 1 (VILIP-1) is a neuronal calcium-sensor protein and a member of the visinin-like protein subfamily [106]. Its physiological role is to regulate neuronal growth,

survival, and synaptic plasticity [107]. Disturbance of calcium homeostasis in neurons, that occurs in AD, causes degeneration of vulnerable neurons and release of VILIP-1 into the extracellular fluid [108]. Because of this characteristic, VILIP-1 has been rated as a marker of neuronal injury. Other biomarkers based on the same pathophysiological process are stanniocalcin-1 [109] and pre-synaptic vesicle protein synaptotagmin [110]. Many studies have confirmed that CSF VILIP-1 is significantly increased in both AD and AD-MCI patients compared to controls and therefore VILIP-1 represents a good diagnostic and prognostic biomarker [111] but unlike Nfl, the correlation between plasma and CSF levels is poor [108].

2.4 Markers of synaptic dysfunction

Synaptic loss and dysfunction are considered to be one of the earliest signs of neurodegeneration [112]. This has led to a high interest in researching synaptic proteins as potential biomarkers that could be useful in diagnosis, prognosis, and guiding treatment of neurodegenerative diseases. Synaptic proteins were first detected in CSF almost 30 years ago [113] and since then the methods for their detection and quantification have advanced significantly, thus increasing their biomarker potential. Today, markers of synaptic dysfunction are mainly focused on one postsynaptic protein, neurogranin, and three presynaptic markers, synaptosomal-associated protein 25 (SNAP-25), synaptotagmin-1, and growth-associated protein 43 (GAP-43) [114].

Neurogranin (Ng), a small calmodulin-binding protein, is expressed mainly by pyramidal cells in the hippocampus and cortex, where it forms granule-like structures [115]. This postsynaptic protein regulates calcium influx via calmodulin and mediates the plasticity, regeneration of synapses, and long-term potentiation [116]. Despite being a good general marker of neurodegeneration, Ng has actually been shown to be more specific for AD than for other neurodegenerative diseases [117,118]. In CSF samples it is possible to detect both full-length Ng and its fragments, mostly short C-terminal peptide species [119]. The function of these Ng fragments is still unknown. However, the evidence suggests that these different fragments have similar predictive value regarding AD [120]. Overall, the research so far points to elevated CSF Ng levels in patients diagnosed with AD, even in prodromal phase of the disease [119,121–133]. Also, it has been shown that CSF Ng levels could be used to predict the progress from MCI to AD [119] and distinct typical from atypical AD forms [134]. A recent meta-analysis [135] confirmed the potential of CSF Ng in predicating memory and executive function decline in subjects diagnosed with MCI. The study also suggests the potential use of CSF Ng/A β_{42} as cognitive function biomarker [135]. These results point to CSF Ng as an useful biomarker for detecting neurodegeneration in the early stages of AD and for differentiating AD from several other AD tauopathies. However, it should be kept in mind that there are also data that do not support the specificity of Ng in AD diagnosis [136].

Synaptosomal-Associated Protein, 25kDa (SNAP-25) is a presynaptic protein which is, along with the vesicle-associated membrane proteins (VAMPs) and syntaxins, a component of the SNARE (soluble N-ethylmaleimide-sensitive factor activating protein receptor) complex, thus playing a role in vesicle formation and neurotransmitter release during synaptic transmission [137]. This presynaptic protein is important in neuronal survival, in the process of neurite outgrowth and long-term potentiation (LTP) [138]. Studies focused on the role of SNAP-25 in AD confirmed higher CSF SNAP-25 levels in AD patients compared to healthy controls, and have demonstrated that these changes can even be detected in the early stages of the disease [75,122–124,139–145]. A recent meta-analysis confirmed that increased CSF SNAP-25 levels can be used to differentiate AD and/or MCI patients from healthy control subjects, which

confirms the potential of this biomarker in the early diagnosis of AD [125,139]. However, elevated CSF levels of SNAP-25 have also been detected in patients diagnosed with other neurodegenerative disorders, including PD, CJD, HD, and it has been associated with psychiatric conditions, including attention deficiency hyperactivity disorder, schizophrenia, and bipolar disorder [146]. The other two presynaptic markers, GAP-43 and synaptotagmin-1, have been less investigated as potential CSF biomarkers in AD. Nevertheless, the level of GAP-43 has been found to be increased in preclinical AD [147–149]. The above discussed synaptic biomarkers were found to have a good discriminating power when trying to distinguish patients with AD from the ones with non-AD dementia [124]. The highest discriminating power for distinguish patients with AD from neurologic controls was suggested for the soluble form of SNAP-25 (SNAP-25aa40) [124].

In addition to the previously mentioned reliable biomarkers of synaptic impairment, other markers of synaptic dysfunction should also be considered in the future studies. One of these markers is a member of the epithelial growth factor (EGF) family, neuregulin 1 (NRG1). NRG1 is involved in neural development, migration and survival of neurons, axon pathfinding, development of glia cells, myelination, and synaptogenesis [150,151]. Proteolytic processing of NRG1 leads to the formation and secretion of soluble forms which interact with post-synaptic receptor tyrosine-protein kinase erbB4 (ErbB4). The levels of NRG1 and ErbB4 were found to be altered in hippocampus and cortex of subjects diagnosed with AD [152,153]. Recent study has shown increased CSF NRG1 levels in subjects with AD and MCI-AD, in comparison to healthy controls and other non-AD dementias [154]. The CSF levels of NRG1 also positively correlated with CSF A β ₄₂, and negatively with MMSE scores [154]. However, the results showed that CSF NRG1 levels, which were found increased in AD and MCI-AD subject compared to controls, had lower discriminatory power than A β ₄₂, t-tau, and p-tau [154]. These results lead to conclusion that markers of synaptic loss and dysfunction poses a great promise as biological measures that could be useful in diagnostics and in therapeutic successes monitoring.

2.5 Markers of neuroinflammation

The presence of neuroinflammation has been well recognized as a concurrent pathological condition in AD. The existence of localized low-level inflammation in the initial stages of AD has been firmly confirmed [155]. Neuroinflammation and cerebrovascular dysfunction are the first occurrences that manifest during the presymptomatic phases of AD and have a role in the further development of the disease [67,83,156]. Both microglia and astrocytes are essential for the initiation and regulation of neuroinflammation [157]. Activated microglia are present in the vicinity of amyloid plaques and play a role in the generation of neurotoxic substances that accelerate neuronal harm [158]. Pathogenic stimuli also trigger astrocytes in AD. Astrocytes have a role in neuroinflammation through the release of pro-inflammatory cytokines, reactive oxygen species, and the facilitation of blood-brain barrier dysfunction [159]. Consequently, this process intensifies the damage to neurons.

The glycoprotein YKL-40, often referred to as Chitinase-3-like protein-1 (CHI3L1), is classified as a member of the chitinase family, although it lacks chitinase activity [160]. In the central nervous system, activated astrocytes and microglia are the main source of YKL-40, which they secrete in response to different inflammatory stimuli and neurodegeneration [161,162]. YKL-40 is involved in several biological processes, including inflammation, extracellular matrix remodeling, cell proliferation, and tissue healing [163]. However, its precise biological role remains incompletely elucidated. The role of YKL-40 in the development of AD and in the

disease's progression is still unclear, but it may be useful as a biomarker for long-lasting brain diseases that have an inflammatory background [164]. People with neurodegenerative diseases have elevated YKL-40 CSF levels, prompting research into its potential involvement in neuroinflammation and neuronal injury [156,165]. YKL-40 has been investigated as a potential therapeutic target, considering its role in various physiological processes and pathological disorders. Modifying its activity or expression may offer promise for treating inflammation, tissue remodeling, and related disorders. A meta-analysis demonstrated that individuals with AD exhibit higher levels of YKL-40 in CSF and plasma compared to healthy controls [166]. These data also suggest a noteworthy association between elevated levels of YKL-40 in CSF and AD. An additional cohort study encompassing 288 people, including both healthy controls and patients diagnosed with various kinds of dementia, assessed the amounts of CSF YKL-40 [167]. Compared to controls, CJD and AD showed higher YKL-40 levels. However, these levels were not significantly higher in VaD or DLB [167]. Wang et al. studied how APOE ϵ 4 affected the levels of YKL-40 in CSF in subjects who were cognitively normal, or were diagnosed with MCI or AD [168]. APOE ϵ 4 carriers had higher levels of CSF YKL-40 than noncarriers with MCI [167]. CSF tau and p-tau concentrations significantly correlated with CSF YKL-40 concentrations in the MCI group [167]. These results show that APOE ϵ 4 may be associated with the amount of CSF YKL-40 in MCI subjects [168]. Another study found an increase in CSF YKL-40 levels in individuals with MCI compared to controls [166]. However, the area under the curve (AUC) was smaller, indicating that YKL-40 has only a modest potential as a biomarker in the context of AD. In a recent study, Abu-Rumeileh et al. discovered that YKL-40 had a moderate diagnostic value, with a sensitivity and specificity of 80% or higher, when comparing controls to AD [169]. A novel study showed that reactive astrocyte biomarkers, like YKL-40, contribute to the impairment of neuronal function and cognitive impairment [170].

Microglia in the CNS predominantly express the cell surface receptor Triggering receptor expressed on myeloid cells 2 (TREM2) [171]. It is a member of the immunoglobulin superfamily and has a vital function in controlling innate immune responses and phagocytosis [171]. The role of TREM2 in microglial activation and response to A β pathology has attracted major attention in the area of AD research. Microglia activation leads to the cleavage and subsequent generation of soluble TREM2 (sTREM2), a measurable indicator of microglial activity in the CSF [172]. Previous research has shown a link between AD pathology and elevated CSF sTREM2, suggesting a strong diagnostic capability [173,174]. The increase in CSF sTREM2 occurs before symptoms appear, but after amyloidosis and neuronal damage [175]. According to a recent study, CSF sTREM2 levels drop in the presence of A β pathology but not tau-related neurodegeneration [175,176]. Just the opposite, the levels of CSF sTREM2 rise in the interaction with tau-related neurodegeneration [175,176]. A notable elevation in CSF sTREM2 levels in non-AD neurodegenerative disorders was found [177]. However, each neurodegenerative disorder has distinct pathological and clinical characteristics. Monitoring TREM2 and YKL-40 may facilitate the assessment of microglial activation and its involvement in the pathogenesis of AD. Moreover, it is important to acknowledge that these indicators possess the capacity to serve as targets for treatment strategies aimed at modulating microglial activation. The therapeutic implications of targeting microglial activation through TREM2 and YKL-40 modulation require further research.

Glial Fibrillary Acidic Protein (GFAP) is a protein that is fundamental in the formation and maintenance of the cytoskeleton of glial cells, specifically astrocytes. It is a reliable marker for reactive astrogliosis and a biomarker for neurodegenerative disorders [178,179]. GFAP can serve as a potent biomarker for predicting dementia risk over a decade before clinical

diagnosis [180]. Elevated CSF GFAP initial levels are indicative of accelerated cognitive decline and associated with significant alterations in other AD biomarkers [181,182]. Given its close proximity to the brain, CSF GFAP is considered to be a precise indicator of brain pathological processes [183]. In the hippocampus of AD patients, especially those with the APOE $\epsilon 4/\epsilon 4$ genotype, GFAP levels are significantly elevated [184]. Greater accumulation of tau in the lateral temporal and frontal areas of the brain was found to strongly correlate with higher levels of GFAP in the periphery [185]. Blood GFAP levels were shown to exhibit a robust correlation with amyloid pathology, making it a more reliable indicator than CSF GFAP [186–188], particularly in predicting the transition from MCI to dementia [189–191]. It is more appropriate to use GFAP as an initial screening strategy rather than a final diagnostic indicator, as it is not exclusive to AD and is implicated also in other neurological disorders [192–194]. Higher levels of GFAP generally correlate with faster declines in cognitive function and contribute to the connection between amyloid pathology and tau protein buildup, ultimately leading to cognitive decline.

2.6 Markers of α -synuclein pathology

Alpha-synuclein (α -syn) is a small (140 amino acids) ubiquitously expressed protein, predominantly found in presynaptic sites in the central and peripheral nervous system [195]. It is encoded by *SNCA* gene and normally, it can exist in two forms, as unfolded monomer or as a folded tetramer of about 58 KDa [196]. It participates in the regulation of synaptic vesicle pool and trafficking [197] and has an important role in assembly of exocytosis mediating SNARE complex [198]. Not only it can form a broad range of structures and associate with lipid and protein chaperones, but under certain circumstances α -syn folds and aggregates into pathogenic forms comprising oligomers, protofibrils, fibrils, which further bring to the formation of protein inclusions [199]. Some post-translational modifications, including phosphorylation, can promote α -syn folding and aggregation which are critical steps in synucleinopathies. Namely, under physiological conditions, less than 5% of monomeric α -syn is phosphorylated, while in pathological protein aggregations such as Lewy bodies, approximately 90% of α -syn is phosphorylated. In typical synucleinopathies, PD, DLB, and multiple system atrophy, neuronal loss is accompanied by the presence of α -syn inclusions. In AD there is an overlap of α -syn, A β plaques, and tau tangle pathologies [200].

Accumulating evidence imply reduced levels of α -syn in CSF of patients with typical synucleinopathies, PD and DLB [201], but a lack of association between PD severity and CSF levels [202]. In a study by Lilamand et al. [203] α -syn levels were found to be significantly lower in CSF of patients with DLB than in patients with AD and authors suggested that CSF α -syn evaluation could improve the early differentiation between DLB and AD. There are also studies showing increased CSF α -syn levels in patients with MCI and AD [204–206], with positive correlation detected between α -syn levels and disease progression [201] or severity of cognitive decline [204,205]. More precisely, CSF α -syn levels were found to be significantly higher in patients with AD with all positive CSF triple markers (A β 42, total tau, and phosphorylated tau) [205,207]. However, there are also opposite results indicating decreased levels of CSF total- α -syn not only in patients with PD and DLB, but also in AD patients compared to healthy control subjects [208]. Some studies demonstrated associations of CSF α -syn concentrations with brain A β deposition measures as well as with CSF t-tau and p-tau181 concentrations [209,210]. This can be further explained by functional studies indicating that α -syn interacts with AD-related proteins. According to results of *in vitro* studies, the interaction of A β with α -syn can accelerate the fibril formation by increasing the aggregation

rate of α -syn [211]. Also, α -syn oligomers can generate and stabilize A β oligomers, leading to fibril-like conformations [212]. It was also shown that α -syn induces tau aggregation, while tau accelerates the fibrillization of α -syn [213]. A study dealing with several plasma and CSF biomarkers in different neurodegenerative disorders, reported significantly higher CSF α -syn levels in patients with AD and MCI in comparison to respective control subjects [214]. However, no significant difference in plasma α -syn levels was found among the same groups of subjects [214].

The α -syn levels are not associated only with AD-related proteins, but also with AD-related genes, such as those coding for presenilin 1 (*PSEN1*) and apolipoprotein E (*APOE*). A recent study found significantly increased tau/ α -syn ratio in AD when compared with healthy controls [207]. Additionally, the ratio was significantly higher in early than in late onset AD [207]. There are reports about the dose-response relationship between the AD risk increasing allele, *APOE* ϵ 4, and CSF α -syn levels, with *APOE* ϵ 4 homozygotes having the highest CSF α -syn levels [204]. Lewy body pathology in the amygdala was reported in the carriers of *PSEN1* mutation among AD patients [215]. Moreover, the frequency of Lewy body deposition was higher in the cases with mutations in *PSEN1* than in those with mutations in *PSEN2* [215]. Although not completely straightforward, these findings imply the important role of α -syn in AD pathology which can be reflected in α -syn measurable manifestations at the periphery. Certainly, a complex interplay between numerous biological processes leading to synucleinopathies still has to be investigated, but results of the studies so far support the idea of α -syn as a biomarker that at least could add to the sensitivity and specificity of standard AD biomarker panel.

3 Conclusions

Biomarkers that have entered routine clinical use (A β ₄₂, t-tau, p-tau181, t-tau/A β ₄₂ ratio), along with other markers that show great potential to be included in the practice itself, are of the great importance in AD diagnosis, not only because they can help distinguish AD from healthy controls, but also due to their ability to help differentiate between different neurodegenerative disorders. In conclusion, CSF biomarkers provide significant additional value in the AD assessment by offering more precise, timely and differential clinical diagnosis of AD at different stages and across different ages. In future, the role of CSF biomarkers in AD will be even more prominent in guiding targeted therapeutic interventions and tailoring patient individual management and support in order to improve the quality of life of both AD patients and their caregivers.

4 Expert Opinion

Considering the direct interaction of CSF with the brain, CSF biomarkers closely reflect the pathophysiological alterations occurring in AD brain. Therefore, it is not surprising that, in addition to neuropsychological evaluation, core CSF biomarkers (A β ₄₂, t-tau, and p-tau181) have been recommended for improvement of timely, accurate and differential diagnosis of AD, as well as to assess the risk and rate of disease progression [20]. Specifically, decreased A β ₄₂ and elevated p-tau levels in CSF suggest A β and tau neuropathology of AD, while increased total-tau concentrations represent a non-specific marker of injured neurons.

There are various advantages of CSF biomarkers. In contrast to PET imaging biomarkers, CSF biomarkers are much cheaper, and could be quickly and simply obtained in a clinical setting [216]. In addition, CSF may enable detection of some biomarkers, which could not be

identified by brain imaging; whereas some CSF biomarker alternations may precede PET biomarker changes; such as elevation of p-tau during AD progression and related cognitive decline [20,83]. However, CSF biomarkers are unable to reflect regional differences in the brain neuropathology that may be particularly important during early AD [83].

Moreover, studies demonstrated that CSF biomarker ratios ($A\beta_{42}/A\beta_{40}$, p-tau/ $A\beta_{42}$, t-tau/ $A\beta_{42}$) could perform even better than individually measured values and may correct for inter-individual differences. In that way, CSF AD biomarker ratios may add relevant information in the differential diagnosis of neurodegenerative diseases, as well as predict the risk of progression from MCI to AD [217].

On the other hand, use of CSF testing may be limited due to perceived invasive nature of lumbar puncture (LB), although several large multicenter studies demonstrated that this procedure is easy, safe and tolerable, with very rare occurrence of serious complications. Further limitations to the routine use of CSF biomarkers include a lack of skills and training in LB procedure, the inability to collect samples from large populations, complex interpretation of the test results, and still present skepticism about their clinical value.

Nevertheless, advanced detection technologies, uniform protocols and standards, as well as fully automated testing procedures that can measure multiple CSF biomarkers in the same sample are now available [20]. Therefore, CSF biomarkers hold promise for a more personalized medicine approach in staging, tracking, and categorization of AD, as well as for the assessing the effects of potential therapeutics [83].

More precise and personalized AD diagnosis may result in lower care costs, delayed institutionalization and reduced mortality, and could be useful for selection of patients suitable to receive novel disease-modifying therapies (DMTs), such as immunotherapies (aducanumab, lecanemab, donanemab) targeting aggregated forms of $A\beta$ [218]. Specifically, via CSF analysis these anti- $A\beta$ mAbs have been confirmed to affect both $A\beta$ plaques, as well as t-tau and p-tau levels [219].

In addition to the core CSF biomarkers, various other CSF markers related to synaptic dysfunction, neuroinflammation, and glial activation (neurogranin, SNAP-25, NfI, YKL-40, TREM2) are now investigated and have yet to be validated for future potential clinical use in early and differential diagnosis, as well as prognosis of AD [20,83].

Moreover, although intensively investigated blood-based biomarkers, as well as biomarkers measured in other fluids, offer relatively non-invasive approach, which is cost-effective and simple to carry, further studies are needed to establish if their clinical utility in AD is comparable to CSF biomarkers [216].

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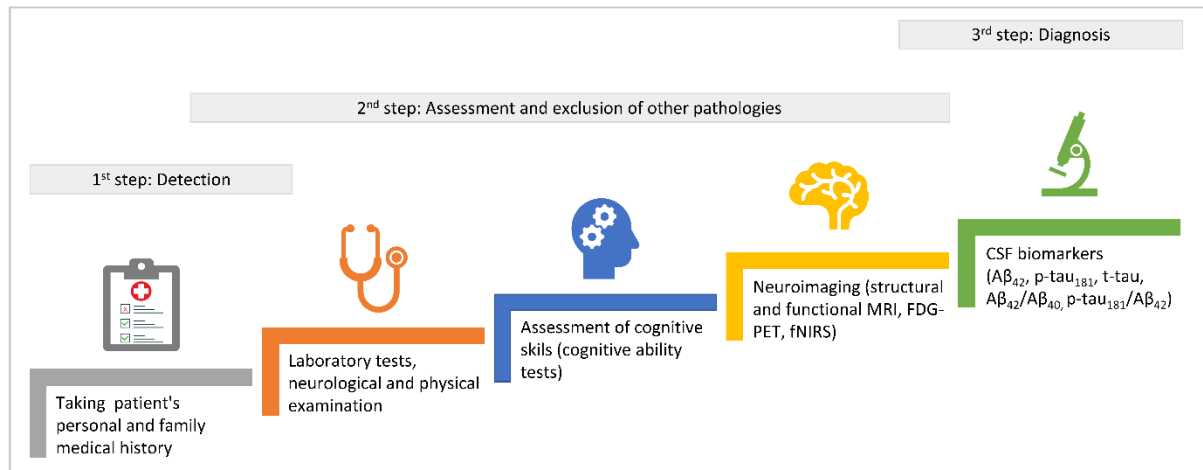


Figure 1. Diagnostic Process in Alzheimer's Disease.

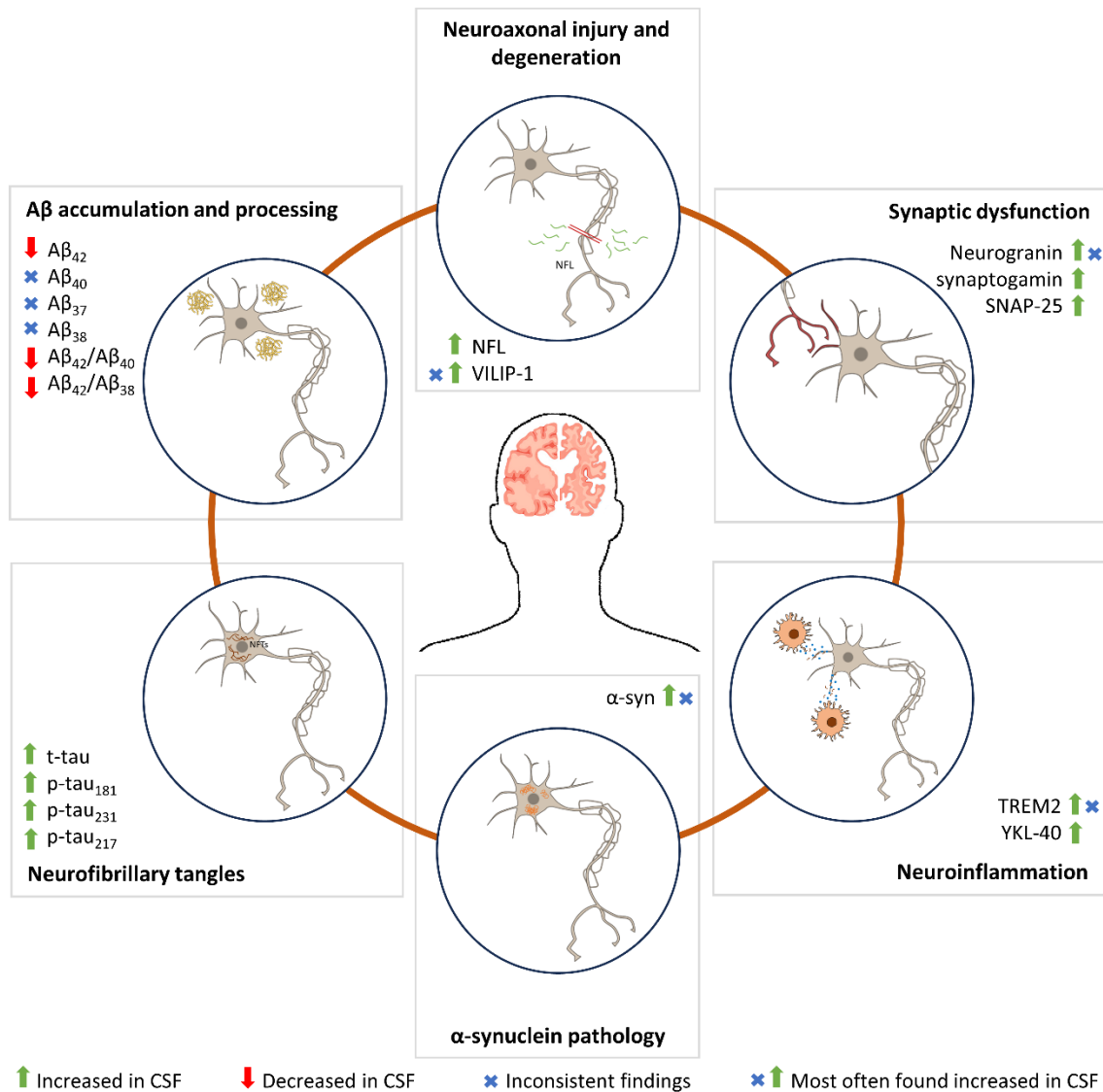


Figure 2. Overview of the most promising CSF biomarkers for Alzheimer's disease diagnosis. Aβ, Amyloid-beta; α-syn, Alpha-synuclein; NFL, Neurofilament light protein; SNAP-25; Synaptosomal-Associated Protein, 25kDa; TREM2, Triggering receptor expressed on myeloid cells 2; VILIP-1, Visinin-like protein 1; YKL-40, Chitinase-3-like protein-1.