



Promising Blood Biomarkers for Clinical Use in Alzheimer's Disease: A Focused Update

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Alzheimer's disease (AD) is the most-common cause of neurodegenerative dementia, and it is characterized by abnormal amyloid and tau accumulation, which indicates neurodegeneration. AD has mostly been diagnosed clinically. However, ligand-specific positron emission tomography (PET) imaging, such as amyloid PET, and cerebrospinal fluid (CSF) biomarkers are needed to accurately diagnose AD, since they supplement the shortcomings of clinical diagnoses. Using biomarkers that represent the pathology of AD is essential (particularly when disease-modifying treatment is available) to identify the corresponding pathology of targeted therapy and for monitoring the treatment response. Although imaging and CSF biomarkers are useful, their widespread use is restricted by their high cost and the discomfort during the lumbar puncture, respectively. Recent advances in AD blood biomarkers shed light on their future use for clinical purposes. The amyloid β ($A\beta$)₄₂/ $A\beta$ ₄₀ ratio and the concentrations of phosphorylated tau at threonine 181 and at threonine 217, and of neurofilament light in the blood were found to represent the pathology of $A\beta$, tau, and neurodegeneration in the brain when using automatic electrochemiluminescence technologies, single-molecule arrays, immunoprecipitation coupled with mass spectrometry, etc. These blood biomarkers are imminently expected to be incorporated into clinical practice to predict, diagnose, and determine the stage of AD. In this review we focus on advancements in the measurement technologies for blood biomarkers and the promising biomarkers that are approaching clinical application. We also discuss the current limitations, the needed further investigations, and the perspectives on their use.

Keywords Alzheimer's disease; biomarker; blood; diagnosis; precision.

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INTRODUCTION

The blood biomarkers that reliably represent Alzheimer's disease (AD)-specific and clinically relevant pathology would aid the care of the patients and the person at risk through easy access to the biomarkers. However, diagnosing and predicting AD using blood biomarkers has been challenging. The difficult journey of proteins, originating in the brain, in systemic circulation across the lymphatic system and blood-brain barrier means that protein levels are much lower in the blood than in the brain. This makes protein for a cerebrospinal fluid (CSF) biomarker difficult to detect using conventional tools in blood. Easy degradation by proteolytic enzymes and nonspecific binding to plasma proteins hampers accurate measurement.¹ The existence of a peripheral source of candidate biomarkers further complicates the matter.²⁻⁴ These hurdles have been overcome with the development of technology that permits measuring proteins at very low concentrations. A series of well-designed studies using an ultrasensitive measurement system elucidated the value of selected blood biomarkers with reference to well-known CSF and neuroimaging biomarkers. The research framework of the ATN system, which sets AD stages based on amyloid β ($A\beta$)

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(A)-, tau (T)- and neurodegeneration (N)-related biomarkers,⁵ has been increasingly applied to the design and interpretation of drugs during their development. The system improves the reliability of a study and increases the chance of detecting a significant finding.⁶ The recent accelerated approval of aducanumab is an example. The positive result on the surrogate endpoint, a reduction of amyloid plaque on amyloid positron emission tomography (PET), was the main basis for approval.^{7,8} However, the accessibility of established biomarkers is restricted by the high cost of PET and the discomfort associated with lumbar puncture. Developing reliable blood-based biomarkers has therefore long been a major objective in AD research, and many efforts have been made. Some blood biomarkers have proven valuable, and these are expected to be added to the ATN framework^{9,10} and clinical practice guidelines.

NEW TECHNOLOGIES FOR MEASURING BLOOD BIOMARKERS

Knowledge of the delicate measurement technologies is indispensable to grasp the recent advances in AD blood biomarkers. Protein biomarker measurements in biological fluids largely depend on immunoassay- and mass spectrometry (MS)-based techniques. Immunoassays require a specific an-

tibody to detect the target protein, while MS measures the target protein by analyzing peptide fragments. The enzyme-linked immunosorbent assay (ELISA) is the most widely used immunoassay in biomarker studies.¹¹⁻¹³ However, its measurement range cannot fully cover the concentrations of blood proteins that are about 2%–10% of those in CSF, when we consider the A β and tau proteins as examples.¹⁴ This may be why previous studies that used ELISA for blood-sample analysis have produced controversial results.¹⁵ Recent progress in technologies has enormously contributed to the establishment of blood-based biomarkers for AD (Fig. 1). Automatic electrochemiluminescence (ECL) technology is increasingly replacing the conventional ELISA to improve test precision even for CSF biomarkers.^{16,17} The Elecsys immunoassay (Roche Diagnostics, Penzberg, Germany),^{18,19} EUROIMMUN ECL immunoassay (EUROIMMUN, Lübeck, Germany),²⁰ and Mesoscale Discovery (MSD) platform (Meso Scale Diagnostics, Rockville, MD, USA)²¹ have been increasingly used to assess blood biomarkers. These fully automatic ECL-based assays root in the sandwich principle that uses two specified antibodies for the capture of respective analytes (antigens) and detection of antigen-antibody complexes. The capture antibody attached to the surface of the working electrode or magnetic beads binds to the target protein after the incubation with the samples. The detector

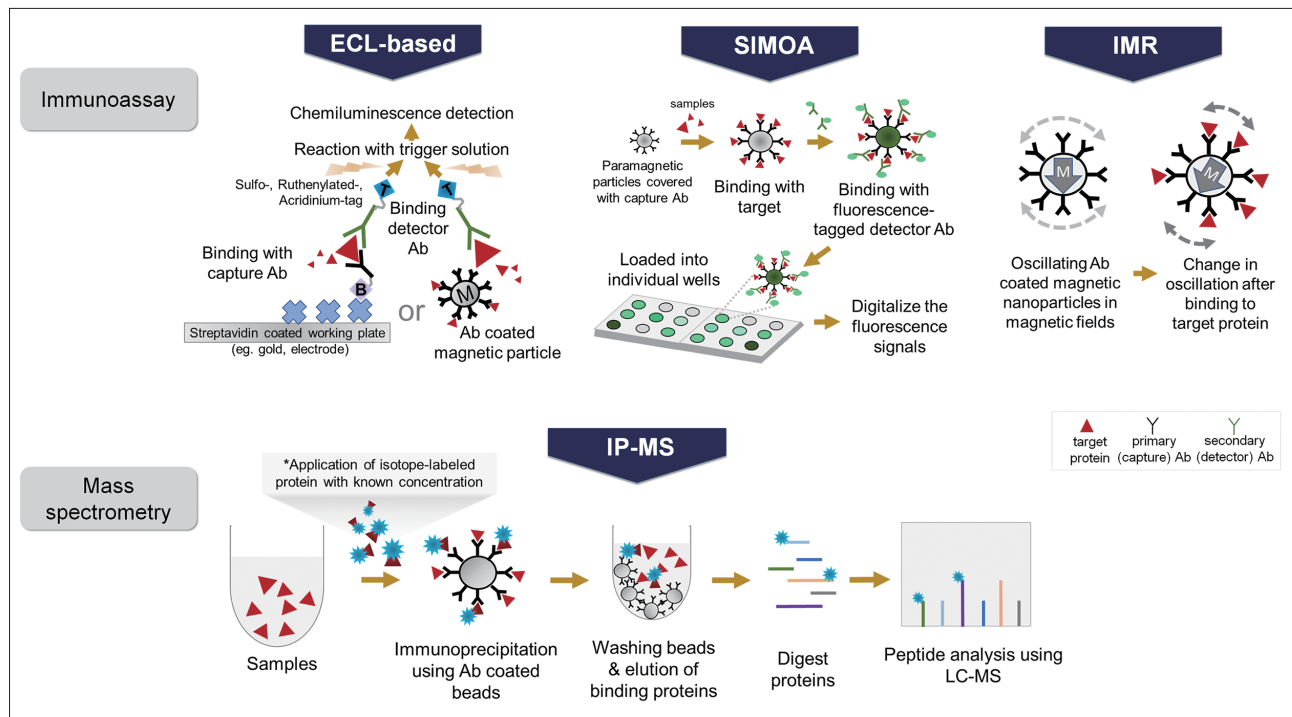


Fig. 1. Ultrasensitive techniques for measuring blood biomarkers. *Additional step for stable isotope spiking absolute quantitation²⁵ to improve amyloid β (A β) peptide quantification using isotope-labeled A β peptide at a specified concentration. Ab, antibody; B, biotin; ECL, electrochemiluminescence; IMR, immunomagnetic reduction; IP-MS, immunoprecipitation coupled with mass spectrometry; LC, liquid chromatography; M, magnetic; MS, mass spectrometry; SIMOA, single-molecule array; T, tag.

antibody is labeled with tags to magnify the signals (e.g., sulfo, ruthenylated, and acridinium tags), and additionally interacts with the target. While being exposed to electrical currents or magnetic fields, the sandwich complexes can remain on the working plate or magnetic beads while the unbound ones are washed out. The complexes are then detected using chemiluminescent signals when the tags of the detector antibodies are exposed to the trigger solution. The strength of the emission is measured and then converted into the target protein concentration.

A single-molecule array (SIMOA; Quanterix, Billerica, MA, USA)¹⁸ and immunomagnetic reduction (IMR; MagQu, Taiwan)²¹ are other types of ultrasensitive immunoassays. The sensitivity of SIMOA is more than 100-fold higher than that of ELISA and more than 25-fold higher than that of the manually performed primitive method of ECL immunoassay¹⁸ by detecting individual paramagnetic beads in a femto-liter-sized chamber, which generates a fluorescent product after binding to the target protein using a coated capture antibody and detector antibody. Since SIMOA digitizes individual fluorophores, the difference between a single immunocomplex can be counted on the beads.¹⁸ The detection accuracy of ECL-based immunoassays is now comparable to or slightly higher than that of SIMOA after automated systems were developed and sensitive antibodies were used on a working electrode such as the MSD platform (<https://www.mesoscale.com/>), as shown in a direct comparative study that measured plasma A β 42/A β 40.¹⁹

IMR has the unique characteristic of using a single antibody and detecting the slowness of rotation of antibody-coated magnetic nanoparticles in a magnetic field after binding to the target antigen. IMR-based quantification of plasma A β 40, A β 42, and total tau (tTau) was recently approved in Taiwan to diagnose and evaluate the risk of AD.²²

MS is a powerful and accurate method for measuring proteins at very low concentrations. Proteomic approaches, which are unbiased and targeted, have also been used to discover and validate AD biomarkers; however, they have been primarily used on CSF samples.²³ Highly accurate MS technology has been applied to detect target proteins at low blood concentrations, such as immunoprecipitation (IP) coupled with MS (IP-MS)²⁴ and IP-MS with Stable Isotope Spike Absolute Quantitation (SISAQ)²⁵ to improve A β peptide quantification while reducing the matrix effect of abundant plasma proteins by adding isotope-labeled A β peptides at a specified concentration. Measuring A β using IP-MS on the SISAQ platform has gained Clinical Laboratory Improvement Amendments and 'CE' approval in the USA and the EU, respectively, surpassing the quality standards for clinical in vitro diagnostic tests.²⁶

Other than the ultrasensitive technique of the conventional ELISA that uses epitope-overlapping A β antibodies to capture and detect A β multimers after spiking the plasma samples, using synthetic A β 42 to enhance the oligomerization of A β ²⁷⁻²⁹ was recently approved by the Korean National Evidence-based Healthcare Collaborating Agency as a supplementary diagnostic tool for AD (#HTA-2021-81, <https://nhta.neca.re.kr/nhta/publication/nhtaU0601L.ecg>).

The approval of more blood-based diagnostic platforms for clinical use is expected to continue in the AD field. However, it should be noted that a biomarker being accepted does not mean that it performs perfectly as an AD biomarker. A large cohort study has started to directly compare the various ultrasensitive techniques to quantify target biomarkers (Table 1)^{19,30} A head-to-head comparison was conducted on the efficacy of eight promising measurement tools (two ECL-based immunoassays, two SIMOA methods, one antibody-free liquid chromatography-MS, and three IP-MS techniques) regarding the degree to which the plasma A β 42/A β 40 ratio reflects the CSF A β 42/A β 40 ratio and positive results in amyloid PET.¹⁹ IP-MS with SISAQ developed at Washington University (St Louis, MO, USA) had the greatest accuracy, with an area under the receiver operating characteristic curve (AUC) of up to 0.872, with low intra- and intertest variabilities.¹⁹ The Elecsys immunoassay from Roche Diagnostics had the best AUC (0.795) and lowest intertest variability among the immunoassays.¹⁹ IP-MS had an AUC (0.817) superior to that of SIMOA (0.620) for both measuring the A β 42/A β 40 ratio and identifying positive amyloid PET findings in subjects without dementia.³⁰ Accuracy, convenience, and cost will determine which technique and target biomarkers will become the most widely used in clinical applications.

A β BIOMARKERS

A β pathology is important for the confirmatory diagnosis of AD.³¹ It is indicated by reduced A β 42 levels in the CSF^{11,32} and increased cortical uptake in amyloid PET.³³ Since these A β biomarkers have been demonstrated to be useful in detecting AD from its early stage, measuring A β in blood has received considerable attention with the aim of establishing AD blood biomarkers. The possible usefulness of the plasma A β 42/A β 40 ratio in predicting and/or diagnosing AD has been suggested through using conventional immunoassay, ELISA, and multiplex techniques, but the results have been inconsistent.³⁴ One of the reasons for this was that a clinical diagnosis that did not consider AD pathology was used for defining the subjects in many studies, although there were several exceptions.³⁵⁻³⁷ This approach does not differentiate subclinical AD from a normal control,³⁸ or non-AD demen-

Table 1. Head-to-head comparison of technologies in measuring the plasma amyloid β ($A\beta$) $_{42}/A\beta_{40}$ ratio

	Elecsys	EUROIMMUN ECL-I	SIMOA N4PE kit	SIMOA Neuro 3 kit	LC-MS-Ar ⁷⁷	IP-MS-WU ^{25,40}	IP-MS-GU ¹⁹	IP-MS-Sh ²⁴
BioFINDER cohort [$n=286$: $A\beta^+/A\beta^- = 118/168$ in CSF, $A\beta^+/A\beta^- = 110/176$ in amyloid PET] ¹⁹								
CSF*	0.778 (0.725–0.832)	0.697 (0.635–0.758)	0.687 (0.626–0.748)	-	0.776 (0.721–0.830)	0.855 (0.810–0.899) [†]	-	-
PET [†]	0.727 (0.669–0.784)	0.672 (0.609–0.735)	0.655 (0.591–0.719)	-	0.753 (0.696–0.811)	0.833 (0.787–0.879) [†]	-	-
BioFINDER subcohort [$n=200$: $A\beta^+/A\beta^- = 86/114$ in CSF and amyloid PET] ¹⁹								
CSF*	0.773 (0.709–0.837)	0.704 (0.631–0.777)	0.679 (0.605–0.753)	-	0.775 (0.711–0.839)	0.872 (0.824–0.920) [†]	-	0.825 (0.767–0.882)
PET [†]	0.773 (0.709–0.837)	0.704 (0.631–0.777)	0.679 (0.605–0.753)	-	0.775 (0.711–0.839)	0.872 (0.824–0.920) [†]	-	0.825 (0.767–0.882)
BioFINDER subcohort [$n=227$: $A\beta^+/A\beta^- = 91/136$ in CSF, $A\beta^+/A\beta^- = 86/141$ in amyloid PET] ¹⁹								
CSF*	0.795 (0.738–0.853)	0.697 (0.628–0.767)	0.706 (0.639–0.773)	0.636 (0.563–0.709)	0.763 (0.700–0.827)	0.838 (0.785–0.891) [†]	0.678 (0.605–0.750)	-
PET [†]	0.728 (0.663–0.793)	0.667 (0.596–0.738)	0.649 (0.577–0.721)	0.600 (0.525–0.675)	0.742 (0.676–0.809)	0.814 (0.760–0.868) [†]	0.632 (0.557–0.707)	-
ADNI cohort [$n=122$: $A\beta^+/A\beta^- = 59/63$ in amyloid PET] ¹⁹								
PET [†]	0.740 (0.651–0.829)	-	0.685 (0.590–0.781)	0.634 (0.534–0.734)	-	0.845 (0.772–0.917) [†]	0.662 (0.565–0.758)	0.821 (0.747–0.895)
					SIMOA, $A\beta_{40}$, $A\beta_{42}$ kits			
					IP-MS³⁰			
British birth cohort [$n=441$: $A\beta^+/A\beta^- = 82/359$ in amyloid PET] ³⁰								
PET [†]	0.620 (0.548–0.691)					0.817 (0.770–0.864) [†]		

Data are the areas under the receiver operating characteristic curves for detecting $A\beta$ pathology, with the 95% confidence interval in parentheses.

*To detect CSF $A\beta$ pathology; [†]To detect amyloid PET positivity; [†]Indicates the highest values.

ADNI, Alzheimer Disease Neuroimaging Initiative; CSF, cerebrospinal fluid; ECL-I, electrochemiluminescence immunoassay; IP-MS, immunoprecipitation coupled with mass spectrometry; IP-MS-GU, IP-MS by the University of Gothenburg; IP-MS-Sh, IP-MS by Shimadzu; IP-MS-WU, IP-MS with SISAQ (Stable Isotope Spike Absolute Quantitation) by Washington University; LC-MS-Ar, liquid chromatography coupled with MS by Araclon; SIMOA, single-molecule array; PET, positron-emission tomography.

tia from AD.³⁹ Another explanation for the discrepancies is the insufficiencies of conventional immunoassays in measuring low blood concentrations of $A\beta_{42}$ and $A\beta_{40}$.

The recent use of fully automated and/or ultrasensitive methods, such as ECL-based assays, SIMOA, and IP-MS technology, with reference to CSF and imaging biomarkers in well-defined subjects has clearly revealed that a decrease in the plasma $A\beta_{42}/A\beta_{40}$ ratio is an indicator of brain amyloidosis.^{19,24,25,40,41} A strong correlation of it with amyloid PET,^{24,41} and with the CSF $A\beta_{42}/A\beta_{40}$ ratio^{19,24,25,40} is observed. In contrast, the plasma $A\beta_{42}/A\beta_{40}$ ratio in nanoparticle-based IMR assays has been found by higher in AD than in controls.^{42,43} This contradiction of IMR was not clearly explained, but it was speculated that IMR increased $A\beta_{42}$ levels by diminishing the binding of $A\beta_{42}$ to plasma proteins in the assay.⁴⁴ An association of increased $A\beta$ level with an AD diagnosis and amyloid PET positivity was also found when using an ELISA-based multimer detection system (MDS) that measures oligomerized $A\beta$.^{27–29} This MDS utilizes the aggregation tendency of blood $A\beta$ to measure multimeric $A\beta$ levels.

The distinctive assay principle of MDS is thought to produce the opposite result.²⁷

The plasma $A\beta_{42}/A\beta_{40}$ ratio has been shown to be valuable in the early detection of amyloid pathology during the early preclinical stage of AD. IP-MS technology seems to be the most-accurate analytical tool for determining that ratio at current. The requirements of a specialized technique and a long processing time, however, is a hurdle for the widespread use of IP-MS.

TAU BIOMARKERS

In addition to $A\beta$ tau deposits forming neurofibrillary tangles in neurons define AD from other neurodegenerative disorders.³¹ CSF τ Tau levels are considered to constitute a neurodegenerative marker, while phosphorylated tau is thought to represent the AD-specific pathology of neurofibrillary tangles.⁴³ Trials to develop CSF-matching blood biomarkers for tau pathology have been performed based on advanced ultrasensitive assays. A series of reports suggested that plasma

phosphorylated tau at threonine 181 (pTau181) and threonine 217 (pTau217) are useful AD biomarkers. An increase in plasma pTau181 levels based on the ECL-based technique using the MSD⁴⁵ and SIMOA⁴⁶ platforms predicts AD development at the preclinical and prodromal stages. An increase in pTau181 level can distinguish AD from non-AD neurodegenerative disorders.⁴⁵⁻⁴⁷ pTau217 is another protein shown to be useful for early detection^{19,48} and for differentiating AD from non-AD dementia.^{47,49} The quantities of pTau181 and pTau217 in plasma are correlated with their corresponding levels in the CSF⁴ and with accumulations of the amyloid and tau proteins in the brain on PET imaging.⁴⁹ These findings collectively suggest that the phosphorylation of these epitopes is AD-specific.⁴ The pTau217 concentration in plasma was found to increase earlier than that of pTau181 according to the amyloid pathology when the two were measured using IP-MS.⁴ The increase in pTau217 preceded the apparent tau accumulation on tau PET,⁵⁰ which suggests that it is potentially useful as the earliest pathology biomarker for tau. The superiority of pTau217 over pTau181 was also identified in the stronger correlation with tau burdens on tau PET,⁴⁷ and the better performance in discriminating AD from non-AD neurodegenerative disorders,^{47,49} monitoring clinical progress,⁴⁸ and predicting dementia onset at the preclinical and prodromal stages.⁵¹

Another plasma form of pTau, which is phosphorylated at threonine 231, was recently measured using the SIMOA platform and was found to be highly accurate in differentiating amyloid PET positivity from negativity and AD from non-AD.⁵² The change in the level of this pTau protein was evident from a very early stage, even before the standard thresholds of amyloid PET positivity and rise in plasma pTau181 and CSF pTau217 levels.⁵² However, a recent direct comparisons of SIMOA-measured pTau181 and pTau231, and between MSD-measured pTau181 and pTau217 indicated that MSD-measured pTau217 followed by MSD-measured pTau181 performed better in predicting abnormal tau accumulation on tau PET at the asymptomatic stage of AD; however, pTau231 revealed a lower performance.⁵³ The contradictory results regarding pTau231^{52,53} increase the need for more investigations into its value as an AD biomarker. Plasma pTau181 and pTau217 also need further validation, particularly considering racial differences. The accuracy of pTau in diagnosing autopsy-confirmed AD varies according to the racial background when assessed using the MSD assay platform. Accuracy was high in non-Hispanic blacks (AUC=0.94 and 0.96 for pTau181 and pTau217, respectively) but low in non-Hispanic whites (AUC=0.65 and 0.75, respectively).⁵⁴ A direct comparison between A β and tau biomarkers in early AD diagnosis has just started. The accuracy of plasma A β 42/A β 40

in IP-MS (AUC=0.817) was higher than that of the pTau181 level (AUC=0.707) on the SIMOA platform in identifying amyloid PET positivity among elderly subjects without dementia.³⁰ However, its accuracy was higher than that of the A β 42/A β 40 ratio (AUC=0.620) when these were both computed using SIMOA. The relative benefits and accuracies of A β and tau biomarkers require further investigation, particularly with the inclusion of plasma pTau217 and pTau231.

NEURODEGENERATIVE BIOMARKERS

In addition to A β and pTau accumulation, which form amyloid plaque and neurofibrillary tangles, neuritic plaque is the most clinically relevant AD pathology.³¹ The product of the dystrophic neural process caused by neuronal injury is the main component of neuritic plaque. The cytoskeletal components of neuronal axons are released into the CSF and then into the systemic circulation when axons are damaged.⁵⁵ Specifically, neurofilament light (NFL), the most abundant of the neurofilament proteins, is detectable in blood using ultrasensitive methods. CSF and plasma NFL levels are correlated,⁵⁶ with both increasing in many neurological disorders such as acute brain injury, stroke, demyelinating disorders, and neurodegenerative dementia.^{55,57,58} Blood NFL levels increase with deterioration of the clinical status and accurately reflect clinical profiles.^{15,59} Tracing the change in serum NFL concentrations has been shown to allow clinicians to predict the onset of dementia 16.2⁶⁰ and 15 years⁶¹ in advance for early-onset AD with a causative mutation. However, the cross-sectional level of serum NFL is less powerful for the early detection of mutation carriers, as this becomes clear only 6.8 years before the expected clinical onset.⁶⁰ The increase in plasma NFL in sporadic AD cases is very delayed, as it is only evident at the prodromal stage (mild cognitive impairment)^{59,62} or after dementia onset.⁵⁷ This is explained by the finding that NFL levels increase with age,^{57,63} and most sporadic AD cases develop at an older age. These results collectively make it more difficult to detect the increase in NFL in response to AD in the elderly population until the symptomatic stage.⁵⁷ The increase in NFL does not persist until an advanced stage of AD. No additional increase in the NFL level or a decrease in the A β 42/A β 40 ratio was observed at a Clinical Dementia Rating (CDR)>1 when the A β 42/A β 40 ratio, pTau181, and NFL in plasma were compared using SIMOA. pTau181 continually increases in AD until CDR=3,⁶⁴ and so clinicians should therefore remain cautious when interpreting NFL levels in subjects at advanced stages of dementia. Blood NFL levels are dynamic, representing the speed of neurodegeneration. They change most dramatically during the rapidly progressive stage⁵⁹ and around the onset of

dementia.⁶⁰

Other than NFL, several axon- and synapse-related proteins in blood have been examined as potential biomarkers that typify neurodegeneration,⁶⁵ but none of them have approached clinical usefulness. The tTau level in CSF is useful as a neurodegeneration biomarker for AD to be incorporated into ATN staging.³¹ However, its plasma levels had a weak correlation with matching CSF tTau levels when measured using SIMOA.⁶⁶ Elevated plasma tTau was suggested to predict the development of all-cause dementia including AD through a meta-analysis⁶⁷ and large cohort study,⁶⁶ however, unlike for NFL, its predictive value was not confirmed in recent studies.^{68,69}

Direct comparisons further indicate the superiority of NFL over tTau regarding the discrimination of AD from healthy controls (AUC=0.94 vs. 0.56;⁶⁸ AUC=0.83 vs. 0.80⁶⁹) and AD from MCI (AUC=0.78 vs. 0.72⁶⁹) on the SIMOA platform. Increased NFL is not an AD-specific event but rather a universal marker of neurodegeneration, independent of causing it. NFL levels are therefore suitable as a screening tool to determine whether a neurological symptom originates

from neuronal damage, and for monitoring the progress and speed of neurodegeneration. Knowledge of the longitudinal change in the level of NFL in the blood instead of its value at a single time point, as well as consideration of AD-specific biomarkers such as Aβ42/Aβ40 and pTau, would simultaneously increase its usefulness as a biomarker.

CURRENT LIMITATIONS AND PERSPECTIVES

Accumulating evidence of the diagnostic and disease-tracking value of blood biomarkers in AD indicates that they will soon be suitable for use. However, several remaining hurdles must be overcome (Fig. 2). First, a consensus on which technique is suitable for the standardized measurements of blood biomarkers in clinical practice is required. Second, reliable cutoff values for individual biomarkers must be determined. Third, a large biomarker gray zone in which differentiation is unclear is problematic when interpreting the available data.^{19,54,70} Fourth, a method for weighting the impact of confounding factors such as aging, comorbidities, and race dif-

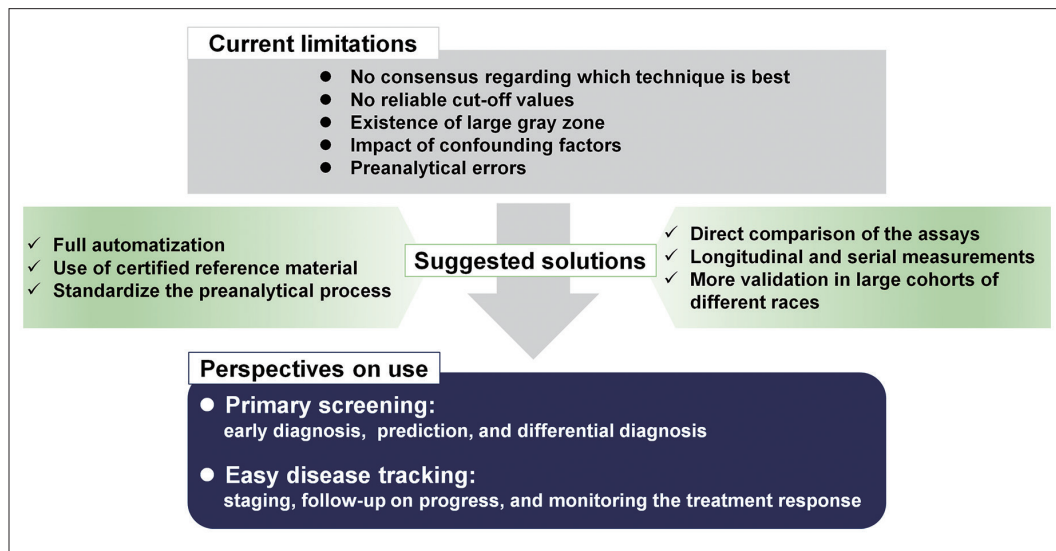


Fig. 2. Current limitations, suggested solutions, and possible clinical applications of blood biomarkers.

Table 2. Blood biomarkers with solid evidence of utility in AD

Biomarker	Early detection	Specific to AD	Correlation with clinical data	Correlation with CSF*	Correlation with PET
Aβ42/Aβ40	Yes, preclinical	Yes	Yes	Yes	Amyloid
pTau181	Yes, preclinical	Yes	Yes	Yes	Amyloid & tau
pTau217	Yes, preclinical	Yes	Yes	Yes	Amyloid & tau
NFL	Yes, prodromal	No	Yes	Yes	Amyloid (asym), tau (sym) [†]

*Correlation with the corresponding level in CSF; [†]Stage-dependent correlation: preclinical asym correlation with amyloid PET, and clinical sym correlation with tau PET.⁷⁸

AD, Alzheimer's disease; asym, asymptomatic stage; CSF, cerebrospinal fluid; NFL, neurofilament light; PET, positron emission tomography; pTau181, phosphorylated tau at threonine 181; pTau217, phosphorylated tau at threonine 217; sym, symptomatic.

ferences on biomarker levels needs to be developed.^{54,71-73} Fifth, the preanalytical errors that profoundly affect assay results must be addressed.⁶¹

The assays should be automated to secure the validity of the biomarkers and corresponding assay tools. The adoption of certified reference materials is necessary to control the assay quality and permit data comparisons between different measurement techniques. Blood-sample handling and the measurement process must be standardized to improve precision. Direct comparing various tools in the same cohort and performing longitudinal tracking instead of single measurements for validation would overcome some of the issues that need to be addressed. Further studies in large community-based cohorts with different racial backgrounds are necessary to establish the utility of individual biomarkers for use on specified technology. The development of certified reference materials for CSF A β 42 measurements according to the Global Biomarker Standardization Consortium⁷⁴ is noteworthy progress in terms of a worldwide trial to monitor assay quality and to harmonize different techniques, with the aim of obtaining reliable cutoff values for fluid biomarkers. Cooperative investigations are expected to continue to improve the precision and utility of biomarkers.⁷⁴

CONCLUSIONS

The accumulating evidence supports the utility of blood AD biomarkers for predicting, diagnosing, and staging AD (Table 2). Identification of AD biomarkers has been achievable thanks to improvements in measurement tools and validation of the results by referring to established CSF and neuroimaging biomarkers. Continuous endeavors to improve and standardize the methods are underway to overcome the current limitations. Blood biomarkers are expected to be used in clinical practice soon. The A β 42/A β 40 ratio, the pTau181 and pTau217, and NFL levels in blood correspond to the pathology of A β , neurofibrillary tangles, and neurodegeneration, respectively. Simultaneously considering these blood AD biomarkers may be helpful in estimating the stage and in differentiating the diagnosis of AD from other brain diseases.⁵ Combining biomarkers with other available data would improve the accuracy of AD diagnoses, as for plasma pTau217 when a brief cognitive test and apolipoprotein E (APOE) genotyping were combined.⁵¹ The most likely scenario is that blood biomarkers will be used for initial screening before more-confirmatory tests, such as CSF biomarkers and PET imaging using a specific ligand, and for monitoring the disease onset, progress, and response to therapy.^{75,76} Adopting blood biomarkers in clinical practice will reduce the economic burden of AD and increase the early and accurate detec-

tion of subjects who will potentially benefit from preventive and disease-modifying treatments when these become available. A focused review of blood biomarkers with solid evidence is timely, and it may help clinicians to prepare for using these biomarkers in medical practice.

Availability of Data and Material

The datasets generated or analyzed during the study are available from the corresponding author on reasonable request.

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Conflicts of Interest

The authors have no potential conflicts of interest to disclose.

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