# A meta-analysis method for identifying potential blood based protein markers associated with Alzheimer's disease

Nivithaa Subburaj<sup>1</sup>, Rani P<sup>1\*</sup> {nivithaasubburaj@gmail.com<sup>1</sup> rani.bio@psgtech.ac.in<sup>1</sup> }

Department of Biotechnology, PSG College of Technology, Coimbatore, Tamil Nadu, India 641004<sup>1</sup>

Abstract. The Alzheimer's disease (AD) is a persistent neurodegenerative disorder characterized by loss of memory and different cognitive capabilities which represents 60 to 80% of dementia. Present day Alzheimer's remedies just treat and postpone the deterioration of dementia manifestations. A biomarker could provide a detailed measurements of brain abnormality, which can aid in the early detection of disease in people who have very mild symptoms and also the treatments targets could be developed for the disease in its early stages, before irreversible brain damage or mental decline occurs. Individuals with Alzheimer's disease and different types of dementia progress at various rates, and biomarkers may aid in predicting and monitoring their progression. Thereby researchers are looking for precise preclinical biomarkers for prognosis of cognitive impairment. The current methods that are used in AD diagnosis are expensive, invasive or time consuming. Hence there is a potential requirement for less invasive and economically feasible blood-based biomarkers to assist in large-scale screening of the geriatric population. The objective of this analysis was to evaluate the reported bloodbased protein biomarkers of Alzheimer's disease (AD). A methodical PubMed survey was performed on 1195 articles distributed among 2007 and March 2018 using the key phrases "Alzheimer's infection" and "plasma biomarker" and 58 articles were chosen that met with the filtering criteria. The blood-based proteins of AD from the selected papers were scored using a meta-scoring system. In this study, 90 blood based proteins were identified, of which 15 were reported multiple times. The six highest meta scored proteins are APOE, BNP, CRP, CD40, TNF a and Clusterin. Further examinations and broad trial validations are important to affirm the clinical use of these potential biomarkers for AD diagnosis.

Keywords: APOE, BNP, CRP, CD40, TNF.

# **1** Introduction

Alzheimer's disease (AD) is a chronic brain disorder that gradually destroys thinking abilities and memory, as well as the ability to perform simple tasks.AD represents about 60% to 70% of dementia cases. Despite the fact that the rate of progression varies, the average life span is around three to nine years once after diagnosis. Dementia affects at least 44 million people worldwide, making it a global health crisis that must be addressed. Around 4 million people are affected from varies forms of dementia in India. Alzheimer's disease affects around 1.6 million people worldwide. In India, the incidence of AD shows elevation in southern part than northern part, whereas comparably lower than China and other parts of the world.

(Mathuranath et al., 2012) Although Alzheimer's affects the population with the age of 65 and above, approximately 200,000 Americans, have earlier onset of Alzheimer's disease based on age.

The cause of Alzheimer's disease is poorly understood. The known risk factors are increasing age, genetics, history of head injuries, depression and hypertension. The amyloid plaques and neurofibrillary tangles in the brain are linked to disease progress in AD. There are no medications or supplements that decrease the risk of acquiring AD. Drugs currently used for AD treatments cannot cure the disease but they can delay the progression of dementia symptoms. Thereby researchers seek highly accurate preclinical biomarkers for early diagnosis in order to minimize cognitive impairment.

Traditional AD biomarker detection has several defects, including invasiveness, low accessibility, high cost, and a narrow scope of clinical applicability. To rule out other possible causes, a probable diagnosis based on the patient's medical records and cognitive assessment (MMSE test), as well as medical imaging and blood tests. Initial symptoms are frequently misinterpreted as signs of normal ageing. Thorough examination of brain tissue for plaques and tangles, is required for a definitive diagnosis. MRI, CT, and PET scans are among the brain imaging techniques used. Blood biomarkers (BBs) have several advantages over CSF and neuroimaging biomarkers, including ease of access and minimal invasiveness.

Despite several attempts to detect AD-related BBs, the selection of reliable BBs has been hampered by low reproducibility. In this study, we will be screening the available protein BBs for AD to identify the prominent biomarkers for clinical diagnosis.

## 2 Methodology

#### A. Article Selection

The 'Alzheimer's Disease' and 'Plasma Biomarker' keywords were used to search the public PubMed electronic database. Out of all the 1195 hits the articles were sorted by publication date from 2007 up till 2018.

B. Exclusion Criteria

Only research articles written in English were considered for the study. Studies were excluded if they looked at other diseases, such as depression and diabetes, or if they didn't provide any clinical data. Studies that dealt with other types of AD biomarkers, such as miRNA, lipids, or ions, or other fields of interest, such as drug efficiency, were also excluded. Only protein biomarkers for which a database exists were chosen for further investigation because the discovery of new biomarkers is dependent on network analysis of previously reported biomarkers. Articles that only report on A $\beta$  and tau proteins were excluded from the meta scoring step because they have been extensively studied previously and found to be less significant for AD diagnosis, and thus their inclusion could introduce bias into the system. Drug trials were excluded because the study's main goal was to find diagnostic markers for Alzheimer's disease. Articles that did not show a consistent change in BB levels were also excluded.

#### C. Inclusion Criteria

All the studies that were considered had to have at least one differentiating feature that distinguished normal subjects from those with Alzheimer's disease. All of the studies used the Mini-Mental State Examination (MMSE) score, which is a widely used scoring system for assessing a patient's cognitive status. Only biomarkers with their own ID were chosen from the

Human Protein Reference Database (HPRD), and biomarkers with the same HPRD ID were counted multiple times.

D. The Meta Scoring System

Each protein BB will be scored using the equation.



S(mi): Meta score	of a biomarker	· mi	C: Set of criterion
mi:Protein	blood	biomarker	S(p, c): Score of a paper with a criterion c
p: Paper			$\mathfrak{S}(p,mi)$ : Significance score of a biomarker
c: Criterion			<i>m</i> i in a paper <i>p</i>

It gives each protein BB a score based on the sum of two multiplied representative values: one is the score of the report stating the protein BB's statistical significance (p-value) derived from that report. Additional scoring was provided if other diagnostic methods were used to identify the AD subjects also: example: MMSE sample + MRI Brain Imaging and if it is a follow up study.

TABLE I Criteria of the meta-scoring system of Blood biomarkers of Alzheimer's Disease

Criteria 1: Number of subjects		Criteria 3: MMSE		
Number of Subjects	Meta Score Given	Welch's t-value of MMSE Range	Meta Score Given	
0	0	NA	0	
0-50	1	<24.15	1	
50-100	2	≥24.15	2	
100-150	3			
>150	4			

Criteria 2: Age		Criteria 4: Statistical Significance		
Welch's t-value of	Meta Score Given	P-Value	Meta Score Given	
age range				
	0	NA	0	
>7 77	1	0.01.0.05	1	
<u>~</u> 2.11	1	0.01-0.05	1	
<2.77	2	0.001-0.01	2	
≥2.77 <2.77	1 2	0.01-0.05 0.001-0.01	1 2	

0-0.001

Table I Indicates four criteria for each report, with a representation of each scoring scale based on its sub-category. Based on their distribution patterns of the number of control and AD subjects scoring scales were assigned

The age criterion is scored by comparing the age of control and AD subjects in each article. The meta-scoring system also used the variation in MMSE scores between control and AD subjects as a criterion. We used Welch's t-test to give the study more weight, despite the fact that studies with greater MMSE differences were believed to have more accurate protein BBs.

To perform Welch's t test, we use the formula

$$t \hspace{0.4cm} = \hspace{0.4cm} rac{X_{1} - X_{2}}{\sqrt{rac{s_{1}^{2}}{N_{1}} \hspace{0.4cm} + \hspace{0.4cm} rac{s_{2}^{2}}{N_{2}}}}$$

Where we compare the AD subjects statistics to the control's statistics (i.e., age and MMSE score) using the standard deviations observed in the study cohorts.

To determine the degree of reliability of each BB, the statistical significance of the protein BBs shown in each article was examined.

## **3 Results And Discussion**

In order to search in the Pubmed database, the keywords such as "alzheimer's disease" and "plasma biomarkers" were used in the study. Out of the 1195 hits in the Pubmed search engine, we had 955 articles upon applying the filtering criteria to exclude review articles, metaanalysis reports, and clinical trials. Reading through the abstracts we have excluded the non human studies, this step had to be done manually as many articles were not Medline indexed. Out of the 955, articles reporting only other types of markers like CSF, MRI, and other Brain imaging techniques to diagnose the AD subjects are excluded. Of the 642 remaining hits, articles reporting only amyloid-beta and tau markers, reports that show no characteristic change in the levels of BB and articles that do not use MMSE scores to measure the level of cognitive impairment (i.e. uses other imaging techniques to confirm AD) were excluded leaving 58 articles which cumulatively report 90 BBs.

Table II summarize the information derived from the articles, while Fig. 1 depicts a schematic diagram of the review process adopted



**Fig. 1**. Flow chart demonstrating the schematic of the systemic review and meta-analysis. Each paper was reviewed and the proteins were listed and meta scored according to the four criterion, extra weightage was given to studies which were a follow up study or if they reported other diagnostic methods also to study the AD subjects. The meta score of all BBs listed were summed up have been reported in Table II

	diagnosis					
No	Biomonitory	Change in biomarker in AD compare	Meta	Count	Defenence	
INO	Blomarker	to control	Score	Count	Keterences	
1	APOE	Increase or	46	6	(Doecke et al., 2012; Guo,	
		decrease			Alexopoulos, Wagenpfeil,	
					Kurz, & Perneczky, 2013;	
					Gupta, 2011; Hu et al.,	
					2012; Llano, Devanarayan,	

TABLE II The list of meta-scored protein biomarkers in blood for Alzheimer's disease

					& Simon, 2013; Soares et al., 2012)
2	BNP	Increase	23	3	(Guo et al., 2013; Hu et al., 2012; Llano et al., 2013)
3	CRP	Decrease	22	3	(Hall et al., 2013; Yarchoan et al., 2013)
4	CD40	Increase	20	3	(Ait-ghezala et al., 2008; Buchhave et al., 2009; Doecke et al., 2012)
5	TNF α	Increase	20	4	(Hall et al., 2013; Huang et al., 2013; Kamer et al., 2009; Kim et al., 2011)
6	Clusterin	Increase	18	3	(Deming et al., 2016; Jongbloed et al., 2015; Schrijvers, Koudstaal, Hofman, & Breteler, 2011)
7	PP	Increase	17	2	(Hu et al., 2012; Soares et al., 2012)
8	sTNFR1	Increase	17	2	(Faria et al., 2014; Zhang, Jia, Qin, & Wang, 2013)
9	SGOT	Increase	16	2	(Guo et al., 2013; Llano et al., 2013)
10	IL 8	Decrease	14	2	(Alsadany, Shehata, Mohamad, & Mahfouz, 2013; Kim et al., 2011)
11	Homocysteine	Increase	13	2	(Doecke et al., 2012; Hall et al., 2013)
12	BACE 1	Increase	12	2	(Goetzl et al., 2015; Wu et al., 2012)
13	BDNF	Decrease	12	2	(Faria et al., 2014; Zheng et al., 2016)
14	MMP 2	Increase	11	2	(Doecke et al., 2012; Lim et al., 2011)
15	A1M	Increase	10	1	(Guo et al., 2013)
16	BIN1	Increase	9	1	(Sun, Tan, Hu, Yu, & Tan, 2013)
17	eotaxin 3	Increase	9	1	(Soares et al., 2012)
18	Gelsolin	Decrease	9	1	(Peng, Jia, & Qin, 2015)
19	IL 16	Increase	9	1	(Guo et al., 2013)
19	sLRP	Decrease	9	1	(Liang, Jia, Wang, Qin, & Liu, 2013)
20	sTNFR2	Increase	9	1	(Zhang et al., 2013)
21	TACE	Increase	9	1	(Zhang et al., 2013)
22	AMP 1	Increase	8	1	(Goetzl et al., 2015)
23	Cathepsin	Increase	8	1	(Goetzl et al., 2015)

24	CysC	Decrease	8	1	(Grewal et al., 2016)
25	DHSM	Decrease	8	1	(Mielke et al., 2011)
26	HNE	Increase	8	1	(Rani, Krishnan, & Rani Cathrine, 2017)
27	IGF II	Decrease	8	1	(Hertze, Nagga, Minthon, & Hansson, 2014)
28	IL 10	Increase	8	1	(Doecke et al., 2012)
29	IL 6	Decrease	8	2	(Huang et al., 2013; Kamer et al., 2009)
30	MAP K	Increase	8	1	(Kiddle et al., 2015)
31	МАРКАРК 5	Increase	8	1	(Kiddle et al., 2015)
32	MMP 3	Increase	8	1	(Peng et al., 2015)
33	MMP 9	Increase	8	1	(Doecke et al., 2012)
34	PAI 1	Increase	8	1	(Oh, Lee, Song, Park, & Kim, 2014)
35	PDS-TTR	Decrease	8	1	(Bradley-Whitman, Abner, Lynn, & Lovell, 2015)
36	sRAGE	Decrease	8	1	(Liang et al., 2013)
37	Tenascin C	Increase	8	1	(Soares et al., 2012)
38	TNF 2	Increase	8	1	(Doecke et al., 2012)
39	TTR	Decrease	8	1	(Velayudhan et al., 2012)
40	Ubiquitin	Increase	8	1	(Goetzl et al., 2015)
41	24S-OH-Chol	Increase	7	1	(Zuliani et al., 2011)
42	ACE	Increase	7	1	(Akatsu et al., 2011)
43	Fibronectin	Increase	7	1	(Lemańska-Perek, Leszek, Krzyanowska-Goląb, Radzik, & Kątnik- Prastowska, 2009)
44	GSH	Decrease	7	1	(Rani et al., 2017)
45	GSSG	Decrease	7	1	(Rani et al., 2017)
46	ICAM 1	Increase	7	1	(Hall et al., 2013)
47	IGBP 3	Decrease	7	1	(Hertze et al., 2014)
48	MDA	Increase	7	1	(Rani et al., 2017)
49	PC Acrolein	Increase	7	1	(Waragai et al., 2012)
50	SM	Decrease	7	1	(Mielke et al., 2011)
51	sNRG -1	Increase	7	1	(Chang et al., 2016)
52	Adiponectin	Increase	6	1	(Une et al., 2011)
53	APO CIII	Decrease	6	1	(Shih et al., 2014)
54	APOA	Decrease	6	1	(Llano et al., 2013)
55	APOE4	Decrease	6	1	(Gupta, 2011)

	1	1		1	1
56	CHI3L1	Increase	6	1	(Choi, Lee, & Suk, 2011)
57	CysC t1	Increase	6	1	(Ghidoni et al., 2010)
58	HCY	Increase	6	1	(Zheng et al., 2016)
59	HDAC	Increase	6	1	(Alsadany et al., 2013)
60	HSP70	Decrease	6	1	(Goetzl et al., 2015)
61	IL 15	Increase	6	1	(Hall et al., 2013)
62	LFP	Increase	6	1	(Chmátalová, Vyhnálek, Laczó, Hort, & Skoumalová, 2016)
63	Neopterin	Increase	6	1	(Parker et al., 2013)
64	NT-Pro BNP	Decrease	6	1	(Marksteiner et al., 2013)
65	PGRN	Increase	6	1	(Piscopo et al., 2013)
66	PON1	Decrease	6	1	(Zengi et al., 2012)
67	VCAM 1	Increase	6	1	(Hall et al., 2013)
68	YKL-40	Increase	6	1	(Craig-Schapiro et al., 2010)
69	YKL-40G	Increase	6	1	(Grewal et al., 2016)
70	APO A1	Decrease	5	1	(Kitamura et al., 2017)
71	C4a	Increase	5	1	(Bennett et al., 2012)
72	CD40L	Increase	5	1	(Ait-ghezala et al., 2008)
73	Fibrinogen	Increase	5	1	(Kitamura et al., 2017)
74	Haptoglobin	Decrease	5	1	(Cocciolo et al., 2012)
75	HO-1	Increase	5	1	(Di Domenico et al., 2012)
76	N-truncated pGlu	Decrease	5	1	(Marcello et al., 2011)
77	SUMO1	Increase	5	1	(Cho et al., 2015)
78	t HODE	Increase	5	1	(Yoshida et al., 2009)
79	t8 iso PGF	Increase	5	1	(Yoshida et al., 2009)
80	α-2 Macroglobulin	Decrease	5	1	(Cocciolo et al., 2012)
81	afamin	Decrease	4	1	(Kitamura et al., 2017)
82	APO A4	Increase	4	1	(Kitamura et al., 2017)
83	BVR-A	Increase	4	1	(Di Domenico et al., 2012)
84	GDNF	Increase	4	1	(Goetzl et al., 2016)
85	IL Iβ	Decrease	4	1	(Kamer et al., 2009)
86	keratin 9	Increase	4	1	(Richens et al., 2016)
87	sAPPα	Increase	4	1	(Goetzl et al., 2016)
88	sAPPβ	Increase	4	1	(Goetzl et al., 2016)
89	α-2-HS-	Decrease	4	1	(Kitamura et al., 2017)
	Glycoprotein				

90 $\gamma$ secretase Increase 4 1 (Goetzl et al., 2016)	4 1 (Goetzl et al., 2016)
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The Protein BB which was reported in most number of articles is APOE with a meta score of 46. Other significant BBs identified in the study are BNP, CRP, CD40, TNF  $\alpha$ , Clusterin, PP, sTNFR1 and SGOT.

The Protein BBs with the highest score are:

A. APOE (meta score=46)

Alzheimer's studies show special curiosity towards ApoE 4 allele, which is capable of enhancing the possibility of late-onset Alzheimer's disease (LOAD) by two- to four-foldwhich is characterized by higher levels of amyloid plaques in the brain. While APOE proteins are higher in the blood plasma and CNS which helps in maintaining the neuronal membrane integrity, the genotype APOE4 is strongly responsible for the lower levels of ApoE protein, marking the onset and progression of AD. The presence of an ApoE 4 allele raised the risk of MCI-related AD progression. However, it's positive and negative predictive values did not support its use as a diagnostic marker forpredicting MCI to AD progression. In the context of research studies, however, it may be useful to enrich or stratify study samples of MCI subjects with faster progression of AD. The number of ApoE 4 alleles has been shown to influence CSF levels of the core AD biomarker candidates A $\beta$ , tau, p-tau, and beta-site amyloid precursor protein-cleaving enzyme-1(BACE1), therefore ApoE genotyping is commonly used in investigatory biomarker studies. The concentrations of serum or plasma ApoE protein levels in Alzheimer's disease have been reported to be inconsistent concluding that the ApoE protein levels cannot currently be recommended as a diagnostic biomarker for AD.(Schneider et al., 2009).

#### *B. BNP* (*meta score*=23)

Gunstad J et al., in 2006 reported a link betweenBNP and cognitive function which was then correlated to the occurrence of new onset dementia in an elderly general population. In accordance with this above association, Hu et al., 2012 and Llano et al., 2013 also reported the elevated levels of BNP in AD.

Also, in the cardiac ventricles, the neurohormone B-type natriuretic peptide is produced. When ventricular myocytes are stretched, preproBNP is released, which is then cleaved into NT-proBNP and BNP by enzymes.Both BNP and NT-proBNP were concluded for their role in diagnosis of heart failure and adding to this, levels of NT-proBNP in plasma is 2-10 times higher than BNP in heart failure patients, with a half-life 3-6 times greater than BNP, suggesting it to be a more appealing biomarker.

C. CRP (meta score=22)

C-reactive protein is produced in the liver in response to inflammation and injury; recent studies correlate the association of CRP levels to AD. According to neuropathology studies, neurofibrillary tangles and senile plaques are linked to CRP in AD brain. A 25-year follow-up study reported that midlife CRP elevations promoted the risk of vascular dementia (VaD) and late-life AD (Schmidt et al., 2002). Once the disease manifests, CRP levels appear to be lower in AD patients in a case-control study.Despite the fact that the significant lower levels of CRP in AD cases has been cross-validated across multiple cohorts, the link has not been validated in different ethnic groups. (O'Bryant et al., 2013)

 $D. CD40(meta \ score=20)$ 

Chronic increased levels of membrane bound CD40 and its ligands CD40L and sCD40L result in the chronic inflammatory response resulting as the continuous activation of microglia in the brain which is strongly associated with AD disease pathology.

In a follow-up study, compared to controls, no significant variation was observed in expression of sCD40 or sCD40L in MCI patients with stable cognition or who hadvascular dementia. But a correlation was seen between sCD40 levels and decreased baseline performance on MMSE. Also a correction was seen between the plasma levels of sCD40 and soluble forms of amyloid precursor proteins (sAPP- $\alpha$  and sAPP- $\beta$ ) in CSF.

sCD40 and sCD40L levels were higher in Alzheimer's disease and they were positively correlated with  $A\beta 1$ –40 and  $A\beta 1$ –42, respectively. They discovered that, combining sCD40, sCD40L,  $A\beta$ , and apolipoprotein E has contributed to high specificity and sensitivity in the diagnosis of AD.

The above studies support the use of sCD40 and sCD40L as AD biomarkers; however, these measurements would be complemented with other biomarkers as a potential candidates with improved specificity and sensitivity. (Giunta, Rezai-Zadeh, & Tan, 2010).

*E. TNF*  $\alpha$  (*meta score*=20)

Chronic inflammation, deposition of amyloid plaques, and intraneuronal accumulation of heavily phosphorylated tau protein forming tangles are the major AD pathological hallmarks. TNF- $\alpha$ , a pro-inflammatory cytokine, is an important molecule in inflammation. Based on genetic and pharmacological manipulations, several *in vivo* studies have shown that TNF-signaling aggravates both A $\beta$  and taurelated pathologies. Both preventive and interventional anti-inflammatory strategies were found to reduce brain pathology and improve cognitive function in AD rodent models (Decourt, Lahiri, &Sabbagh, 2016).

F. Clusterin (meta score=18)

Clusterin is a glycoprotein that primarily serves as an extracellular chaperone. Clusterin variants are strongly associated with the late-onset Alzheimer's disease as shown in large GWAS (Genome-Wide Association Studies), earning clusterin a spot on AlzGene's top 10 risk genes.

Clusterin levels are higher in AD brain, but it's unclear how the protein affects AD pathogenesis. This could be due to the fact the brain's ability to respond to stressors are impaired by pathogenic clusterin variants. The ability of clusterin to bind to A $\beta$  peptides and thus influence their aggregation, deposition, and/or clearance is a popular hypothesis, but the underlying mechanism(s) is still unknown. Clusterin variants are being linked to neurological phenotypes in functional studies. Clusterin can be found in blood and cerebrospinal fluid, but it has not yet proven to be a useful biomarker. (Clusterin | ALZFORUM., 2018)

According to Park et al. (2014), while the number of studies reporting certain high scoring BBs such as APOE, clusterin, and CRP has increased, the number of studies reporting new biomarkers is still limited. Additional research is required even in the case of high meta-scored protein Blood Biomarkers in order to confirm its clinical use as potential biomarkers and toimprove the mode of diagnosis.

## **4** Conclusion

Alzheimer's disease is a chronic brain disorder which ischaracterized by three major groups of symptoms: psychiatric symptoms, behavioral disturbances and cognitive impairment. Alzheimer's disease appears in a variety of forms; vary from mild memory loss to severe dementia. AD patients are frequently identified and treated in primary-care settings, can impose diagnostic and management challenges. Currently, stringent episodic memory tests (MMSE tests) are the only available best predictors of transition from mild cognitive impairment to AD.Cognitive impairments in multiple cognitive arenas are observed several years before the clinical onset of Alzheimer's disease that are stable for several years. (Burns & Iliffe, 2009)

Early detection of cognitive impairment, as well as clinical assessment and management, postpones the need for nursing home care and lowers the risk of false diagnosis and irrelevant treatment strategies. The advantages of early Alzheimer's disease diagnosis include the initiation of symptomatic treatments for minimal cognitive impairment.

Even though imaging techniques can detect early changes in the brain, there is no single technique that could specifically identify people with MCI who can develop AD or other dementias if used as a screening test.

Traditional AD biomarker detection has a number of flaws, including invasiveness, high cost, limited accessibility and a narrow focus on clinical applicability. Blood biomarkers (BBs) have several advantages over CSF and neuroimaging biomarkers, including ease of access and minimal invasiveness.

There is an existing systemic review for identification of protein based blood biomarkers reporting the BBs that were identified from 1989 till march 2013 (Park et al., 2014). The objective of the present study is to review and score the BBs reported currentlythat could potentially be identified as useful in a clinical setting for diagnosis.

The Protein BB which was reported in most number of articles was APOE with a meta score of 46. Other significant BBs identified in the study are BNP, CRP, CD40, TNF  $\alpha$ , Clusterin, PP, sTNFR1 and SGOT. Further studies are necessary even for high meta-scored protein BBs, to validate the potential biomarkers.

It appears unlikely that a peripheral biomarker would replace specific and consistent markers that are closer to the disease state such as PET imaging markers and CSF molecular markers. Peripheral biomarker will be less specific, but potentially more sensitive. They are undoubtedly easier to replicate in a broader population that necessarily involve repeated measurements to monitor disease progression. As an early step toward diagnosis in life, a funnel-based approach of blood-based biomarkers could be used. This would be a significant step toward making clinical trials effective and feasible.

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