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## Blood biomarkers for Alzheimer’s disease in clinical practice and trials

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### Abstract

Blood-based biomarkers hold great promise to revolutionize the diagnostic and prognostic work-up of Alzheimer’s disease (AD) in clinical practice. This is very timely, considering the recent development of anti-amyloid- $\beta$  (A $\beta$ ) immunotherapies. Several assays for measuring phosphorylated tau (p-tau) in plasma exhibit high diagnostic accuracy in distinguishing AD from all other neurodegenerative diseases in patients with cognitive impairment. Prognostic models based on plasma p-tau levels can also predict future development of AD dementia in patients with mild cognitive complaints. The use of such high-performing plasma p-tau assays in the clinical practice of specialist memory clinics would reduce the need for more costly investigations involving cerebrospinal fluid samples or positron emission tomography. Indeed, blood-based biomarkers already facilitate identification of individuals with pre-symptomatic AD in the context

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of clinical trials. Longitudinal measurements of such biomarkers will also improve the detection of relevant disease-modifying effects of new drugs or lifestyle interventions.

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A neuropathological diagnosis of AD is based on the presence of widespread cortical plaques containing A $\beta$  fibrils in combination with neuronal neurofibrillary tangles and neuropil threads containing hyperphosphorylated tau<sup>1</sup>. Tau-containing tangles restricted to the medial temporal lobe are found in most people older than 60 years. In AD, A $\beta$  plaques start to accumulate 10–30 years before dementia onset, and these changes are thought to facilitate the spread of pathological tau species from the medial temporal lobe throughout the neocortex<sup>2</sup>. The mechanism by which A $\beta$  aggregates drive tau spread and accumulation is not yet known but could involve increased tau phosphorylation and secretion of soluble tau forms<sup>3</sup>. Even though tau pathology affects different cortical regions in a rather stereotypic order<sup>2</sup>, there is evidence that spreading of tau might occur along four main trajectories, resulting in four main tau patterns that are associated with somewhat different clinical syndromes and prognoses<sup>4</sup>.

A relatively large number of drugs have been developed against A $\beta$  and tau. Immunotherapies targeting aggregated A $\beta$  have recently been shown to be very effective at removing A $\beta$  fibrils from the brains of patients with AD, which has been associated with beneficial clinical effects<sup>5–7</sup>. For example, lecanemab was recently approved for clinical use in the USA to slow down the clinical deterioration of symptomatic patients with AD<sup>7</sup>. With the introduction of effective disease-modifying therapies in clinical practice, we urgently need scalable and cost-effective methods for accurate diagnosis of patients with early AD. Unfortunately, the diagnostic work-up of AD is rather mediocre when biomarkers are not used to support the clinical diagnosis. In specialized memory clinics, the misdiagnosis of AD is around 25–30% when not using AD-specific biomarkers<sup>2</sup>. However, the vast majority of individuals with AD are managed in primary care where >50% of AD cases are not routinely recognized or correctly diagnosed, resulting in suboptimal treatment and care, which is especially problematic in light of emerging disease-modifying treatments for AD<sup>2</sup>. However, the recent development of blood-based biomarkers (BBMs) for AD holds promise to revolutionize the diagnostic work-up of AD in clinical practice globally but also to improve the design of clinical trials for the earliest stages of AD. In this Review, we will briefly discuss current cerebrospinal fluid (CSF) and imaging biomarkers for AD, which are already used in certain specialist memory clinics. Next, we will discuss recently developed BBMs for AD and how they can be used in both specialist memory clinics and primary care as well as in clinical trials. An overview of current fluid and imaging biomarkers is given in Table 1.

## Current imaging- and CSF-based biomarkers for AD

### Imaging-based biomarkers

There are several positron emission tomography (PET) tracers that can detect the load of A $\beta$  fibrils in the brain. Three A $\beta$  PET tracers (flutemetamol, florbetapir and florbetaben) are approved for clinical use, and several large-scale studies have shown high concordance between the in vivo uptake of these PET tracers and the density of A $\beta$  plaques as determined

post-mortem<sup>2</sup>. A normal A $\beta$  PET scan result rules out AD as the underlying etiology in most patients with cognitive symptoms; an abnormal A $\beta$  PET scan is indicative of AD in a younger patient with cognitive symptoms, but, in an older patient, such a result should be interpreted with caution, considering that about 40% of individuals aged 90 years have A $\beta$  plaques in the brain<sup>8</sup>.

Several PET tracers can detect the load of insoluble tau aggregates in the brain<sup>2</sup>. One tau PET tracer (flortaucipir) is approved for clinical use in the USA. This tracer has been validated against neuropathology, and it can reliably detect the density of both neurofibrillary tangles and neuropil threads<sup>9,10</sup>, although it lacks the sensitivity to reliably detect the earliest tau stages (restricted to the medial temporal lobe)<sup>10</sup>. Tau PET has shown excellent diagnostic accuracy for distinguishing AD dementia from most other neurodegenerative diseases<sup>11</sup>, and it has been suggested that this method can be used to rule in AD in patients with cognitive impairment even at older ages, considering the high specificity of neocortical tau PET retention for patients with AD<sup>12</sup>. In a recent study, cognitively unimpaired individuals with both positive A $\beta$  PET and positive tau PET had 20 $\times$  and 40 $\times$  increased probabilities of developing mild cognitive impairment (MCI) and dementia, respectively, compared to those with normal PET scans<sup>13</sup>. Cognitively normal individuals with positive A $\beta$  PET but negative tau PET had a very minor risk of developing cognitive impairment<sup>13</sup>. Together, these results support the National Institute on Aging-Alzheimer's Association (NIA-AA) research framework for AD, which states that individuals with both A $\beta$  and tau pathology should be labeled as AD independent of cognitive status (that is, including cognitively unimpaired individuals)<sup>14</sup>.

### CSF-based biomarkers

A $\beta$  and tau can also be measured in CSF<sup>2</sup>. CSF A $\beta$ <sub>42</sub> levels and especially the ratios of A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> or A $\beta$ <sub>42</sub>/p-tau correlate strongly with A $\beta$  PET status<sup>15,16</sup> and AD neuropathology<sup>17</sup>. Several CSF A $\beta$  and p-tau assays on high-performing, fully automated platforms are currently used in clinical practice<sup>16,18</sup>. Given the high degree of agreement between A $\beta$  PET and CSF A $\beta$ , there is usually no need to perform both investigations on the same patient<sup>19</sup>.

Tau can be phosphorylated at more than 40 different positions. Tau phosphorylation at threonine 181 (p-tau181) is increased in CSF in AD but not in other neurodegenerative diseases, including other tauopathies<sup>20</sup>. Other p-tau isoforms have also been investigated extensively in CSF, and there is converging evidence that p-tau217 levels exhibit stronger associations with both tau tangle and A $\beta$  plaque load than levels of p-tau181 and p-tau231 (refs. 21,22), although some results indicate that the assay setup may be more important than the phosphorylation site as such<sup>23,24</sup>. Furthermore, CSF p-tau217 levels might distinguish AD dementia from other dementias with even higher accuracy than other p-tau isoforms, and this has improved prognostic utility<sup>21,22,25,26</sup>.

### Comparing PET- and CSF-based A $\beta$ and tau measures

According to the NIA-AA research framework for AD, A $\beta$  pathology (A) can be determined using either A $\beta$  PET or CSF A $\beta$  in an interchangeable fashion<sup>14</sup>. This is likely to be correct

in most situations<sup>27</sup>, but there are subtle differences between these two measures. First, levels of CSF A $\beta$ <sub>42</sub>, and potentially also A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub>, change somewhat earlier than A $\beta$  PET signals; this is also the case for A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> levels in blood samples<sup>28–30</sup>. Furthermore, the A $\beta$  PET signal increases with disease progression as it measures insoluble A $\beta$ -laden plaques, whereas, in CSF and blood, the A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> ratio decreases with development of pathology.

However, using tau PET and CSF p-tau interchangeably for tau pathology (T) seems to be more complex; for example, in cognitively unimpaired populations, more individuals are identified as T-positive when using CSF p-tau versus tau PET<sup>27</sup>. This is because p-tau levels in CSF and plasma start to increase much earlier than the tau PET signal reaches the threshold for detection during the preclinical stages of AD<sup>31,32</sup>. In fact, A $\beta$ -positive individuals who are positive for CSF p-tau but still negative for tau PET might represent a population with early AD who are just about to start accumulating tau aggregates in the neocortex<sup>33</sup>. It has therefore been suggested that the NIA-AA research framework be updated to include p-tau and tau PET as separate biomarker entities, that is, using ‘APT’ instead of ‘AT’, where P stands for p-tau (measuring the levels of soluble hyperphosphorylated tau) and T stands for tau PET (measuring the density of insoluble tau fibrils)<sup>33</sup>.

### Markers of neurodegeneration

Finally, according to the NIA-AA research framework, markers of neurodegeneration (N) provide additional information about disease status<sup>14</sup>. Hippocampal volume and/or cortical thickness of temporoparietal regions can be determined using structural magnetic resonance imaging (MRI) and reflect the disease stage of AD. Furthermore, several fluid biomarkers of neurodegeneration have emerged. For example, CSF levels of total tau (t-tau) reflect axonal degeneration and injury. Disorders with rapid neurodegeneration, such as Creutzfeldt–Jakob disease and autoimmune encephalitis, are characterized by normal CSF p-tau but a more pronounced increase in t-tau<sup>34,35</sup> than that found in AD (which has a slower clinical course). Similarly, in acute neuronal injury such as stroke and acute brain trauma, CSF t-tau shows a temporary increase associated with severity of the neuronal damage and long-term clinical outcome, while p-tau remains relatively normal<sup>36,37</sup>. Another promising neurodegeneration biomarker is neurofilament light (NfL), which reflects axonal degeneration and injury of the longer myelinated axons of the brain and spinal cord structures, irrespective of cause. NfL levels in CSF are especially increased in amyotrophic lateral sclerosis, frontotemporal dementia and atypical parkinsonian disorders but also in AD<sup>38</sup>. Importantly, in most neurodegenerative disorders, higher levels of NfL are associated with faster disease progression and higher brain atrophy rates<sup>38,39</sup>. NfL can therefore be regarded as a measure of the intensity of ongoing neurodegeneration. Even though a substantial number of CSF markers for neurodegeneration and neuroinflammation have been developed over the past decade (Table 1), only A $\beta$ , tau and NfL seem to provide clinically relevant prognostic information in the context of AD<sup>40</sup>.

## BBMs for AD and related disorders

As in CSF, plasma levels of A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> are associated with the presence of A $\beta$  plaques in the brain as determined by neuropathology<sup>41</sup>. In many studies across several platforms, including different immunoassays and mass spectrometry-based assays, the plasma A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> ratio is lower in A $\beta$ -positive groups than in A $\beta$ -negative groups, regardless of cognitive status of the cohort<sup>42–47</sup>. However, the performance of different plasma A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> assays varies substantially, and a recent head-to-head comparison showed that certain mass spectrometry-based assays could detect A $\beta$  pathology with areas under the receiver operating characteristic curve (AUC) of 0.84–0.87, whereas many commonly used immunoassays performed much worse (AUC, 0.64–0.69)<sup>48</sup>. Adding *APOE* genotype to plasma A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub>-based prediction models increases the AUC by about 10% (refs. 45,47,48). The assays with better diagnostic performance are characterized by superior control of measurement error. Still, these relatively high-performing A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> assays exhibit only modest correlations between the levels in plasma and CSF ( $r_s$  of 0.56–0.65)<sup>48</sup>, probably because much of the A $\beta$  in plasma is derived from peripheral sources<sup>49</sup>.

Several high-sensitivity assays have recently been developed that can reliably detect different p-tau isoforms in plasma, including p-tau181 (refs. 50–53), p-tau217 (refs. 54,55) and p-tau231 (ref. 56). These assays performed well in detecting AD as defined using neuropathology<sup>50–54,56</sup>. A few head-to-head comparisons of these assays using plasma from patients with cognitive complaints showed that assays quantifying plasma p-tau217 are somewhat better at detecting AD pathology and predicting future development of AD dementia<sup>57–60</sup>. The best-performing p-tau217 assay showed a high correlation between plasma and CSF levels, with a correlation coefficient of 0.89 (ref. 57). Plasma p-tau231, on the other hand, seems to start increasing at very low A $\beta$  plaque levels<sup>56,61,62</sup>. These results are congruent with recent studies showing that plasma p-tau231 is associated with A $\beta$  plaque load but not tau tangle load<sup>41,63</sup>. By contrast, p-tau181 and p-tau217 were associated with both A $\beta$  plaques and tau tangles, with p-tau217 showing stronger correlations with both pathologies<sup>41,64</sup>. Our current understanding is that A $\beta$  plaques might induce hyperphosphorylation and secretion of tau, which in turn might promote tau aggregation and formation of tau tangles<sup>3,64–66</sup>. However, there is currently no tangle-specific tau plasma marker, but recent developments in CSF markers hold great promise, especially those measuring the microtubule-binding region (MTBR) of tau<sup>67</sup>.

Similar to CSF NfL, plasma NfL is a measure of active neurodegeneration in several neurodegenerative disorders<sup>68</sup>. Plasma NfL levels generally correlate well with the levels in CSF<sup>69</sup>. NfL levels are associated with neurodegeneration in AD, but the effect size is smaller for plasma than for CSF, as is the case in other neurodegenerative diseases, for example, Huntington's disease<sup>70</sup>.

Glial fibrillary acidic protein (GFAP), which probably reflects reactive astrocytes, can be reliably measured in both blood and CSF. Plasma levels of GFAP are increased in individuals with early A $\beta$  pathology<sup>71–73</sup> and can predict subsequent cognitive decline and conversion to AD dementia in cognitively unimpaired individuals<sup>74</sup> and in patients with MCI<sup>75</sup>. Plasma GFAP levels are also increased in other neurodegenerative diseases,

including frontotemporal dementia associated with progranulin mutations<sup>76</sup>. It is currently unclear whether plasma GFAP levels correlate with the number of reactive astrocytes as determined post-mortem using immunohistochemistry or antemortem using PET.

## **BBMs for diagnosis and prognosis of cognitively impaired patients in specialist memory clinics**

### **BBMs as diagnostic biomarkers in clinical practice**

Once anti-A $\beta$  therapies (for example, lecanemab) can be used in patients with MCI or mild dementia, it will be crucial that a highly accurate yet time- and cost-effective diagnostic workflow for AD is in place. BBMs hold great promise in this respect (Fig. 1). In clinics without access to A $\beta$  PET or CSF AD biomarkers, implementation of accurate AD BBMs will improve the diagnostic work-up quite substantially compared to the care as usual of today. In specialist clinics with access to CSF and/or PET, BBMs will speed up the diagnostic process and substantially reduce costs. BBMs will probably be sufficient to support or reject an AD diagnosis in most patients with MCI or dementia; only those patients with uncertain BBM outcomes are likely to need confirmatory testing with A $\beta$  PET or CSF AD biomarkers (Fig. 1). Indeed, a recent study showed that a diagnostic algorithm based on plasma p-tau217 resulted in an accurate AD diagnosis in about 80% of patients with MCI, whereas around 20% had uncertain blood biomarker results and needed further confirmatory testing with CSF AD biomarkers<sup>77</sup>. A newly developed, highly accurate mass spectrometry assay for p-tau217 might result in fewer patients with uncertain biomarker outcomes, reducing the need for CSF and PET even further<sup>57</sup>.

An important question is which plasma biomarkers for AD should be implemented in the assessment of patients with MCI and dementia. Although plasma GFAP and NfL levels are increased in patients with MCI or dementia due to AD, they are unlikely to contribute substantially to accurate detection of AD pathology when combined with high-performing plasma p-tau and A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> assays<sup>78,79</sup>. By contrast, several different p-tau variants, including p-tau181, p-tau217 and p-tau231, are clearly increased in the plasma of patients with MCI or dementia due to AD, and these can be used to distinguish AD from other neurodegenerative diseases with high diagnostic accuracy, often on par with PET and CSF AD biomarkers (for reviews, see, for example, refs. 2,80–82). Plasma p-tau217 is the tau variant that shows the largest fold increase in individuals with symptomatic AD, with increases of about 300–700% compared to both healthy individuals and patients with other neurodegenerative diseases<sup>54</sup>. Therefore, the clinical performance of this biomarker is less susceptible to test–retest variability when compared to many other plasma biomarkers<sup>83</sup>, and the effects of comorbidities (for example, kidney dysfunction) on plasma p-tau217 levels are minor<sup>84</sup> (see below). The latter is especially true when the p-tau217/t-tau217 ratio is used as quantified using mass spectrometry<sup>85</sup>. Together, these characteristics of plasma p-tau217 result in robust clinical performance of this biomarker for detection of AD in patients with MCI or dementia in clinical practice (Fig. 2). However, plasma levels of p-tau217 are very low in healthy individuals, and it might therefore be challenging to establish this biomarker on many of the fully automated platforms used in clinical practice today, as has been the case for the Roche Elecsys platform<sup>79</sup>. Although plasma p-tau217 is currently the

best-performing diagnostic biomarker for symptomatic AD, there are also high-performing assays for plasma p-tau181 (refs. 57,58), and plasma levels of p-tau181 are generally higher than p-tau217 and therefore easier to measure on fully automated platforms<sup>79</sup>.

When it comes to plasma  $A\beta_{42}/A\beta_{40}$  levels, the very modest drop of 8–15% in symptomatic AD<sup>2</sup> means that this biomarker has low performance and robustness in routine clinical settings, even if analytical variability and systematic bias are kept at a minimum<sup>86</sup>, and few current  $A\beta_{42}/A\beta_{40}$  assays fulfill this requirement, resulting in large variability in the clinical performance of different plasma  $A\beta$  assays<sup>48</sup>. Nevertheless, high-performing plasma  $A\beta_{42}/A\beta_{40}$  assays can contribute to plasma p-tau-based diagnostic algorithms that are designed to detect AD pathology in patients with MCI<sup>78</sup>.

A recent consensus paper proposed that high-performing BBMs can already be used in specialist clinics to facilitate detection of AD pathology in patients with MCI or dementia<sup>87</sup>. Importantly, BBMs should be combined with a thorough clinical assessment, including psychiatric and neurological examinations, cognitive testing and structural brain imaging. BBMs should never replace such investigations, and they should only be used in patients with cognitive impairment for whom AD is a possible diagnosis and where such a diagnosis will probably change the management of the patient<sup>87</sup>. These recommendations are primarily based on the risk that false-positive results could lead to anxiety, depression or rash behavior; even a 5% false-positive rate would mean thousands of people would be inappropriately diagnosed with AD if the tests were used in broad screening ahead of identification of objective cognitive impairment.

### **BBMs as prognostic biomarkers in clinical practice**

Information about individual-level prognosis is of key interest for patients with mild cognitive complaints as well as for their care partners and responsible physicians<sup>88</sup>. Higher baseline plasma p-tau217 and p-tau181 levels in patients with mild cognitive complaints are associated with subsequent progression to AD dementia<sup>51,89–91</sup>. Combining continuous values of plasma p-tau217 (or p-tau181) levels with performance on a few brief cognitive tests outperforms predictions made by dementia experts and performs similar to CSF-based prognostic models when predicting development of AD dementia within 2–6 years in patients with mild cognitive complaints<sup>90</sup> (Fig. 2). An easy-to-use online tool based on plasma p-tau217 and three brief cognitive tests can be used to determine the prognosis of individual patients, and similar tools are likely to be used in clinical practice in the near future<sup>90</sup>. Neither plasma NfL nor  $A\beta_{42}/A\beta_{40}$  contributed much when predicting the development of AD dementia<sup>90,91</sup>. However, plasma NfL might have a value when predicting future decline of global cognition in patients with MCI or dementia: prognostic models based on plasma p-tau and NfL can predict changes in global cognition (Mini-Mental State Examination (MMSE) and Clinical Dementia Rating Scale Sum of Boxes (CDR-SB)) in patients with MCI, with performances similar to those of models based on CSF biomarkers<sup>91</sup>. However, a recent study showed that tau PET imaging may have an even greater value for predicting global cognitive decline in patients with MCI or dementia and that plasma NfL is the only plasma biomarker that provides any additional prognostic

information<sup>92</sup>. However, tau PET is costly and currently not widely available in clinical practice.

### **BBMs as prescreening biomarkers in clinical trials**

Plasma AD biomarkers will also facilitate recruitment of patients with MCI or dementia due to AD for clinical trials. About 40–60% of patients with MCI and 20–30% of those with clinically diagnosed AD dementia do not have brain A $\beta$  pathology<sup>2</sup>. Thus, when recruiting patients with prodromal AD or mild AD dementia for clinical trials, prescreening individuals for, for example, plasma p-tau217, would reduce the need for confirmatory investigations involving A $\beta$  PET or CSF AD biomarkers (Fig. 3). Such prescreening with high-performing BBMs is likely to be more cost effective in patients with MCI than in patients with dementia, considering the lower prevalence of A $\beta$  positivity in MCI. In certain interventional AD trials, such as trials evaluating lifestyle interventions, a high-performing plasma biomarker might be enough to confirm AD pathology, removing the need for CSF and PET altogether, which would substantially reduce the costs and increase the scalability of such trials.

## **BBMs for diagnosis and prognosis of cognitively unimpaired individuals in specialist memory clinics**

### **BBMs as diagnostic biomarkers in clinical practice**

High-performing assays for plasma p-tau181, p-tau217, p-tau231, A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> and GFAP but not NfL can detect relatively well AD-related pathological changes in cognitively normal individuals and in patients with subjective cognitive decline (for reviews, see, for example, refs. 2,81,82). A recent study analyzed all these plasma biomarkers in preclinical AD and showed that plasma p-tau231 and A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> could be used to detect the earliest AD brain changes<sup>61</sup>. Indeed, a combination of plasma p-tau and A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> was found to be the best biomarker combination for detection of amyloid pathology in cognitively unimpaired individuals, and high-performing A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> assays might contribute more to the diagnostic work-up in this very early disease stage than to later disease stages<sup>78</sup>. Although there is currently no obvious clinical need to detect AD in cognitively normal individuals, this might change when phase 3 trials evaluating anti-amyloid therapies, such as lecanemab (NCT04468659) and donanemab (NCT05026866), will read out in 2027–2028. That said, the use of BBMs might be considered in certain patients with subjective cognitive decline, for whom cognitive test results are still normal but the patient history indicates a gradual cognitive deterioration. Such patients could be investigated with BBMs in clinical practice similar to patients with MCI (see above).

### **BBMs as prescreening biomarkers in clinical trials**

Even if AD BBMs will not be widely used for cognitively normal individuals in clinical practice in the foreseeable future, they will be a gamechanger for clinical trials conducted in patients with preclinical AD. As only 10–30% of individuals aged 60–80 years are positive for amyloid PET or CSF A $\beta$ <sup>8</sup>, a large number of PET (or CSF) examinations is currently needed to identify a sufficient number of individuals for phase 3 trials focusing



on preclinical AD. For the A4 (Anti-Amyloid Treatment in Asymptomatic Alzheimer's) trial (the first phase 3 trial for preclinical AD), it took 3.5 years and more than 4,000 amyloid PET scans to identify 1,169 participants eligible for the study. As shown in Fig. 3, a prescreening step with high-performing BBMs could greatly reduce the number of PET (or CSF) investigations. Using high-performing plasma  $A\beta_{42}/A\beta_{40}$  assays in this way indeed resulted in substantial cost and time savings<sup>46,93,94</sup>. This was particularly evident if the plasma test was incorporated early in the enrollment process, even before the screening visit<sup>93</sup>. As mentioned above, combining plasma  $A\beta_{42}/A\beta_{40}$  with p-tau231 (ref. 61) (or p-tau217 (ref. 78)) levels might result in even more efficient detection of preclinical AD. Several large-scale phase 3 anti- $A\beta$  trials already use plasma  $A\beta_{42}/A\beta_{40}$  ([NCT05026866](#)) or p-tau217 ([NCT04468659](#)) to identify individuals with a high probability of having preclinical AD.

As shown in Fig. 3, efficient clinical trials also need to enrich the preclinical AD population for those that will probably worsen in the primary outcome over a reasonable time period (3–5 years). This is because many individuals with preclinical AD do not deteriorate over 5–10 years or even during their lifespan<sup>95–97</sup>, and, without enrichment for vulnerable individuals, very large and extended trials would be needed. Power calculations indicate that, if only amyloid positivity is included as a requirement, about 2,000 participants are needed per group to detect a treatment effect of 25% over 4 years using a cognitive composite measure optimized for preclinical AD as a primary endpoint<sup>95</sup>. In two independent cohorts, plasma p-tau217 levels could accurately predict future cognitive decline in preclinical AD; in this setting, plasma p-tau217 performed better than other plasma and CSF biomarkers (p-tau231, p-tau181, GFAP and NfL) or amyloid PET<sup>98</sup>. Importantly, power calculations revealed that using plasma p-tau217 levels to enrich for cognitively normal individuals likely to show cognitive decline resulted in large reductions in required sample sizes.

Tau pathology has consistently been shown to be more strongly associated with clinical deterioration than with  $A\beta$  pathology, even in cognitively unimpaired individuals<sup>13,99</sup>. Therefore, future phase 2 trials might use accumulation of tau pathology over time (as measured with longitudinal tau PET) as a more precise primary outcome than cognitive measures, which exhibit high intra-individual variation. Of note, the increase in tau PET signal over time in amyloid-positive AD populations is modest. However, plasma p-tau217 was recently shown to accurately predict future accumulation of tau aggregates in the brain, and a combination of p-tau217 and tau PET at baseline could be used to substantially reduce the needed sample sizes by >40% when using longitudinal tau PET as the primary outcome in preclinical AD trials<sup>100</sup>.

## Potential use of BBMs in primary care settings

Most patients with cognitive symptoms are managed in primary care rather than in specialist clinics. Although few studies in primary care settings have systematically evaluated the accuracy of AD diagnoses against a valid reference standard (for example, dementia expert diagnoses supported by CSF or PET), it seems that about 50–70% of patients with cognitive impairment are currently not recognized or correctly diagnosed in primary care, due to lack of easily accessible, time- and cost-effective, and accurate diagnostic tools<sup>101</sup>. The problem

is even worse in early stages of the disease, that is, in patients with subjective cognitive decline or MCI, because there are no accurate methods for personalized prognosis of AD in primary care. This leads to patients not receiving appropriate diagnostic and prognostic information and also results in suboptimal treatment strategies and care. Misdiagnosis can also lead to unnecessary care seeking and costly investigations due to diagnostic uncertainty. Considering that CSF and PET cannot be used in primary care, AD BBMs have the potential to finally provide primary care physicians with adequate tools to provide their patients with an accurate diagnostic and prognostic work-up.

Several prospective studies are currently evaluating AD BBMs in primary care. For example, a study in Sweden that includes 800 patients with cognitive symptoms at primary care centers evaluates whether AD BBMs can be analyzed prospectively in primary care using pre-defined cutoffs in a diverse population in which many patients have several comorbidities, whether diagnosis and treatment of patients improve by adding AD BBMs to the 'care as usual' and whether BBMs can be used to predict future development of AD dementia in non-demented individuals with cognitive complaints in primary care. Regulatory authorities in many countries will probably require such studies before AD BBMs can be widely implemented in primary care settings, which is why the Alzheimer's Association appropriate-use recommendations do not yet endorse the use of AD BBMs in primary care<sup>87</sup>. Once BBMs for AD have been validated in primary care, education packages regarding when to use the biomarkers, what they represent, how to interpret the results and what to do with the results must be developed in close collaboration between primary care physicians, dementia experts and patient representatives<sup>87</sup>.

## BBMs for monitoring disease progression

Fluid biomarkers and brain-imaging methods are increasingly being used as outcome measures in clinical trials evaluating disease-modifying therapies for AD and other neurodegenerative disorders. The use of such surrogate endpoints will be especially important in preclinical AD trials, for which very large and long-term studies are needed when using a clinical outcome such as cognitive function<sup>95</sup>. Biomarker outcomes predicting clinical beneficial effects could shorten the duration and/or reduce the size of future preclinical AD trials. A $\beta$  PET but not yet any AD-related fluid biomarker is deemed by the US Food and Drug Administration (FDA) to be a 'reasonably likely surrogate endpoint', which means that it is 'supported by strong mechanistic and/or epidemiologic rationale, but the amount of clinical data available is not sufficient to show that they are a validated surrogate endpoint' (ref. 102). Such a biomarker can be used to support the FDA's Accelerated Approval Program. However, only validated surrogate endpoints can be used as a primary endpoint in pivotal trials used for full FDA approval, and no AD biomarker currently meets this definition.

Many of the fluid tau and neurodegeneration biomarkers discussed above are more or less directly related to disease progression. The best-established biomarker for general neurodegeneration is NfL<sup>68,103</sup>. The magnitude of NfL increases in CSF and/or plasma reflects the intensity of the neurodegenerative process and predicts imaging and clinical evidence of disease progression<sup>104,105</sup>. In AD, high NfL levels are associated with

longitudinal neurodegeneration as determined by MRI; however, this is only obvious at more advanced dementia stages<sup>105</sup>. Such associations are clearer in other neurodegenerative diseases such as multiple sclerosis<sup>106</sup>, amyotrophic lateral sclerosis<sup>107</sup> and frontotemporal dementia<sup>108</sup>, in which NfL levels are generally much higher than in AD<sup>68</sup>. Interestingly, disease-modifying treatment in, for example, multiple sclerosis and spinal muscular atrophy reduces NfL levels, and the reductions correlate with the clinical efficacy of the intervention<sup>109,110</sup>. In anti-A $\beta$  antibody trials for AD, attenuated increases of CSF NfL have been reported<sup>111,112</sup>, but no such results have been obtained thus far for plasma NfL<sup>113</sup>. NfL may be a better surrogate marker for neurodegenerative disease other than AD, considering the modest increases in plasma NfL in AD and considering the fact that many older individuals have other brain pathologies (for example, TDP-43) that are more related to increased NfL levels than AD.

Early studies showed that people with clearly increased CSF tau levels had faster AD progression, suggesting that this marker, similar to NfL, reflects the intensity of the neurodegenerative process in AD<sup>114,115</sup> but in an AD-specific rather than in a general neurodegeneration-reflecting manner. Similarly, studies with new blood tests for p-tau forms<sup>2,80–82</sup> showed that longitudinal changes in plasma p-tau levels are associated with both brain atrophy and cognitive decline in AD populations<sup>116–118</sup>. Importantly, promising anti-A $\beta$  antibody trials have shown treatment-induced reductions in plasma p-tau markers associated with less clinical deterioration, supporting disease modification and slowing of the neurodegenerative process<sup>5,113</sup>. In clinical practice, it is possible that certain plasma p-tau forms will be used to assess the effect of anti-A $\beta$  antibody treatments for both treatment evaluation and disease-monitoring purposes. One could even envision yearly plasma p-tau testing to detect reoccurrence of disease activity, if and when treatment with anti-A $\beta$  antibodies for 1–2 years eventually becomes a reality.

In addition to p-tau and NfL, other markers of disease intensity that predict AD progression and have shown promising results in clinical trials include plasma GFAP. Plasma GFAP levels increase over time in AD<sup>75</sup>, and clear reductions are observed after efficient removal of A $\beta$  plaques by anti-A $\beta$  immunotherapy<sup>113</sup>. Furthermore, CSF and plasma A $\beta_{42}$ /A $\beta_{40}$  ratios have been suggested to detect drug target engagement of anti-A $\beta$  antibodies. However, therapeutic antibodies may change the half-life of the biomarkers, making data interpretation difficult<sup>119</sup>, as has been reported for biofluid-based tau biomarkers in anti-tau antibody trials<sup>120</sup>.

Few longitudinal studies have performed head-to-head comparisons of different plasma AD biomarkers. Recently, we reported that plasma p-tau<sub>217</sub> increases more clearly over 4–6 years in preclinical and prodromal AD than A $\beta_{42}$ /A $\beta_{40}$ , p-tau<sub>181</sub>, p-tau<sub>231</sub>, GFAP and NfL; p-tau<sub>217</sub> also had the strongest associations with brain atrophy and cognitive decline in two independent cohorts<sup>61</sup>. If replicated in other studies, this might indicate that plasma p-tau<sub>217</sub> could be a key biomarker for detecting disease-modifying effects in drug trials and other interventional studies (for example, involving physical activity) targeting preclinical and/or prodromal AD stages.

## Standardization, robustness and clinical cutoffs of BBMs

### Standardization

Before biomarker-based diagnostic tests can be introduced into routine clinical practice, biomarker standardization and the development of certified reference materials and guidelines are essential to assure high quality of laboratory test results (and thereby patient care and safety), specifically the accuracy of diagnostic classifications.

For the core AD CSF biomarkers, a working group under the International Federation of Clinical Chemistry and Laboratory Medicine has led standardization efforts<sup>121</sup>. These have resulted in mass spectrometry methods for CSF A $\beta$ <sub>42</sub> that have been approved by the Joint Committee for Traceability in Laboratory Medicine as reference measurement procedures. They have also resulted in three certified reference materials (low, medium and high A $\beta$ <sub>42</sub> levels) intended to be used to calibrate the commercially available immunoassay, thereby harmonizing levels across assays<sup>122</sup>. Similar standardization efforts have been initiated for AD BBMs. A first round-robin study (which aims to verify a new method and compare results across methods and laboratories) on A $\beta$  methods showed disappointingly poor correlations across plasma A $\beta$ <sub>42</sub> assays ( $r = 0.41$ – $0.54$ ), including mass spectrometry methods, whereas correlations for A $\beta$ <sub>40</sub> assays were better ( $r = 0.59$ – $0.79$ )<sup>123</sup>. Using the A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> ratio did not improve correlations<sup>123</sup>, and another study obtained similar results<sup>48</sup>. When the same immunoassays are applied for CSF samples, correlations are generally very high ( $r = 0.94$ – $0.99$ )<sup>124</sup>. In contrast to plasma A $\beta$ <sub>42</sub>, correlations between different high-performing plasma p-tau assays are tight<sup>57,60</sup>.

A more widespread launch and implementation of the AD blood biomarkers for clinical use will require not only analytical standardization but also ensuring that blood biomarkers can be measured on the type of laboratory analyzers available in non-specialized, smaller hospital laboratories. Methods for the measurement of these AD BBMs on high-precision, fully automated instruments have been published<sup>79</sup>, and other assay formats have been released as laboratory-developed tests for potential clinical implementation.

Standardization of sample-collection procedures is also crucial for clinical implementation. Pre-analytical sample-handling procedures have been examined extensively for CSF biomarkers, as such factors may affect biomarker values<sup>125</sup>. For blood biomarkers, the same type of sampling tubes (EDTA plasma) should be used for all biomarkers; all the blood biomarkers can withstand up to three freeze–thaw cycles<sup>126,127</sup>. In contrast to CSF A $\beta$ , plasma A $\beta$  is not sensitive to collection tubes made of glass, and tubes with a gel separator can be used. Importantly, both A $\beta$ <sub>42</sub> and A $\beta$ <sub>40</sub> are unstable in whole blood, with levels decreasing already after 2 h; samples should therefore be centrifuged early (optimally within 1 h) and plasma should be separated, after which it can be stored at +4 °C for up to 6 h before freezing<sup>126,128</sup>.

### Robustness

Robustness describes a biomarker's ability to classify patients with high consistency and high clinical accuracy<sup>80,83,86</sup>. For a biomarker to be suitable for clinical use, its levels should be clearly higher (or lower) in AD samples than in all relevant differential diagnostic groups,

resulting in high diagnostic sensitivity and specificity (Box 1). The effect size, meaning the difference in mean biomarker levels between patients with and without AD or the pathology (for example, brain amyloidosis) divided by the pooled standard deviation, needs to be much larger than the total measurement variability of the BBM. The total variability depends on biological variability, variability induced by variations in pre-analytical handling of blood samples and the analytical variability inherent to any measurement technique and to drifts or changes over time (Box 1). In other words, a robust biomarker can withstand the variability and bias across measurements that occur in clinical routine. As examples, the pregnancy test for urine human chorionic gonadotropin (HCG) has very high robustness, because levels in pregnant compared with non-pregnant women are more than 1,000-fold different. Also the  $A\beta_{42}/p\text{-tau}$  and  $A\beta_{42}/A\beta_{40}$  ratios are very robust when measured in CSF<sup>129</sup>. By contrast, the  $A\beta_{42}/A\beta_{40}$  ratio has very low robustness when measured in plasma, as the ratio is only 0.9-fold lower in patients with brain amyloidosis than in amyloid PET-negative cognitively normal older people<sup>48</sup>.

Factors that may affect biomarker measurements are shown in Box 1. Factors that contribute to biological variation can influence classification accuracy and may need consideration when establishing cutoffs (see below). In addition, both pre-analytical (for example, time to centrifugation) and analytical (assay imprecision) factors and drifts or bias in values across rounds of measurements will also add to the total measurement variability (Box 1).

A blood biomarker such as the plasma  $A\beta_{42}/A\beta_{40}$  ratio, which exhibits a modest change of 8–15% in amyloid-positive cases, may be problematic even if the total measurement variability is lower than 5–10% (ref. 86). This small effect size, combined with the total error in plasma  $A\beta_{42}$  assays, means that this biomarker has low robustness. This may induce difficulties if the plasma  $A\beta_{42}/A\beta_{40}$  ratio is introduced as a clinical routine test. By contrast, plasma p-tau217 levels are increased 300–700% in symptomatic AD<sup>54</sup>.

Biomarker robustness can be tested through simulations that test the influence of increasing the analytical total error of blood biomarker measurements on clinical classifications. Such simulations have shown that even minor increases in total error strongly affect the performance of the plasma  $A\beta_{42}/A\beta_{40}$  ratio as a biomarker to identify brain  $A\beta$  pathology but not that of other blood biomarkers (NfL, GFAP and p-tau181)<sup>130</sup>. A second study found that introducing a 10% bias had a large effect on performance of the plasma  $A\beta_{42}/A\beta_{40}$  ratio but not the CSF p-tau/ $A\beta_{42}$  ratio when they were used as biomarkers of amyloid positivity<sup>86</sup>. A third study showed that, even though plasma  $A\beta_{42}/A\beta_{40}$  has lower test–retest variability than plasma p-tau217, NfL and GFAP, plasma p-tau217 was least affected by this test–retest variability with a change in diagnostic accuracy of <1% (ref. 83). The better robustness is due to p-tau217 having a substantially higher effect size than the other BBMs<sup>83</sup>. Consequently, plasma p-tau217 and p-tau181 seem to be robust AD BBMs<sup>83,130</sup>. Of note, the robustness might depend on disease stage: the effect size increases with severity of pathology, because there is a gradual increase in fold change from preclinical AD to prodromal AD, with the highest levels in AD dementia<sup>50,51,53,54</sup>. Thus, even if these biomarkers are very robust in symptomatic AD, they might be less robust in detecting preclinical AD, which may have implications for prescreening in preclinical AD trials (see above).

## Clinical cutoffs

Regarding the clinical diagnostic performance of the biomarkers, it should be noted that, in principle, all data published thus far come from retrospective studies, in which all samples were analyzed in batch, after which the optimal cutoff was identified and descriptive data on the performance were calculated (AUC, sensitivity and specificity). To generate data on the 'real-life' diagnostic performance, prospective studies are needed, with fixed biomarker cutoffs set before the start of the study and biomarkers analyzed on a routine (daily or weekly) basis, allowing the obtained biomarker results to be influenced by all the components constituting the true total measurement variability (Box 1).

For use in clinical practice, biomarkers need well-defined and widely accepted clinical cutoffs. Ideally, each biomarker should have a cutoff value established based on the discrimination between clinical groups (or established proxies for neuropathology) or, alternatively (and commonly used in laboratory medicine), based on the 95th percentile of values in a well-characterized control group<sup>131</sup>.

Baseline physiological levels of brain proteins in blood depend on various non-disease-associated factors. For example, blood NfL levels are strongly age dependent<sup>132</sup>. Studies assessing sex differences in blood biomarkers have shown inconsistent results. Although it is common to have age- or sex-specific normative ranges for laboratory tests used in clinical practice, comorbidities are typically left as risks or contraindications to the test or simply need to be considered by the patient's physician during interpretation of the test result. Indeed, several comorbidities (for example, chronic kidney disease and obesity) are associated with increases in plasma p-tau<sup>84</sup> and plasma A $\beta$ <sub>40</sub>, A $\beta$ <sub>42</sub>, NfL and GFAP<sup>133</sup>, even though A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> (ref. 134) and p-tau<sub>217</sub>/t-tau<sub>217</sub> (ref. 85) ratios seem to be unaffected. Of note, although associations between blood biomarkers and comorbidities may be statistically significant in large clinical or population-based cohorts, it is important to describe the magnitude of such effects, especially the effect size<sup>135</sup>, and whether it is of clinical relevance. Indeed, in two large clinical cohorts, plasma NfL and GFAP and, to a lesser degree, p-tau were associated with kidney dysfunction and body mass index, but these potential confounders had no clinically meaningful effects on either prediction of brain pathophysiology or future cognitive change<sup>134</sup>. In line with these results, chronic kidney disease, obesity and other comorbidities affect the reference ranges for the AD blood biomarkers only slightly<sup>84,134</sup>. As mentioned above, biomarker cutoffs in laboratory medicine are not routinely adjusted for comorbid disorders, but it is useful to understand their influence on biomarker results, as they might confound interpretation at an individual patient level (for example, in a patient with severe kidney disease and obesity).

Although common laboratory tests (for example, hemoglobin, platelet count and  $\gamma$ -glutamyl transferase) show differences across racial or ethnic groups<sup>136</sup>, reference intervals for normality are usually developed predominantly with white populations and not separately for different subpopulations. Possible differences in blood biomarker levels across racial or ethnicity groups have also been discussed, but recent large studies on the BBMs A $\beta$ <sub>42</sub>, A $\beta$ <sub>40</sub>, t-tau, p-tau and NfL found that levels were similar across white, Black and Spanish-speaking Americans<sup>137,138</sup>. These results suggest that the same cutoff for AD BBMs can be used across racial or ethnicity groups. However, further studies are needed to assess possible

physiological differences in blood biomarker levels across ethnic groups, also adjusting for socioeconomic status and comorbidities.

When BBM levels are close to the established cutoffs, the interpretation is more uncertain. Patients with such uncertain results could be referred for confirmatory CSF or PET testing (Fig. 1). Indeed, categorization of individuals into low-probability ('non-AD'), high-probability ('AD') and intermediate-probability ('gray zone') groups has been suggested for the most common AD biomarkers, and a combined model using several markers resulted in fewer patients in the intermediate-probability ('gray zone') group<sup>83</sup> (Fig. 1). A similar classification system is used for a test available for clinical use in the US, a probability score based on combining *APOE* genotype, age and plasma  $A\beta_{42}/A\beta_{40}$  ratio<sup>139</sup>. As mentioned above, a p-tau217-based diagnostic algorithm could classify about 80% of patients with MCI correctly as having or not having AD, with 20% ending up in the intermediate-probability ('gray zone') group<sup>77</sup>.

### Future directions

AD is a common disease for which promising drugs are now emerging that may slow or even stop  $A\beta$ -triggered breakdown of neuronal networks. Disease-modifying drugs with different targets (for example, anti-tau therapeutics and synapse stabilizers) are also underway. The emerging availability of this broader range of potentially disease-modifying drug candidates directed against distinct pathogenic mechanisms in the AD process resembles recent developments in, for example, rheumatology, for which effective targeted treatments started to become available 20 years ago and have now been implemented in clinical practice in close collaboration between primary healthcare physicians and specialists using biomarker-supported personalized medicine approaches. We envision similar developments in AD in the next few years, and the recently developed BBMs will play a very important role in this process.

We envision that individuals presenting to primary care physicians with cognitive concerns will be first examined according to standard clinical procedures, starting with an evaluation of the patient's medical history, present comorbidities and duration of cognitive symptoms, a basic neurological examination and brief cognitive testing. The clinician can subsequently make a request for BBM testing after having discussed its potential implications with the patient and his or her relatives. Elevated levels of plasma p-tau would suggest that AD pathology is responsible for the observed cognitive impairment, whereas normal plasma p-tau levels would indicate non-AD causes. If p-tau is normal, increased blood NfL concentration could suggest the presence of non-AD neurodegeneration. We must stress, however, that BBMs might help the clinician in decision making but should in no case substitute a proper neurological assessment. Indeed, confirmatory diagnosis in specialist care settings will continue to be important for some time for many patient populations, but, in the future, it will probably be possible to accurately diagnose and treat many of these patients in primary care only.

The ability of plasma p-tau measurements to identify AD pathophysiology in individuals with symptomatic disease demonstrates the potential of this marker for identifying and recruiting  $A\beta$ -positive symptomatic participants for clinical trials. In addition, we expect

that blood p-tau will be important for the recruitment of pre-symptomatic A $\beta$ -positive cohorts, which will result in reduced rates of negative PET scans and substantial cost and time savings. Plasma p-tau biomarkers will also be useful to evaluate effects of therapeutic intervention: significant decreases in plasma p-tau concentration or a reduction in the rate of increase over time could indicate beneficial effects of anti-A $\beta$  treatments.

The discussions above point to a revolution in the next 2–4 years, in which widespread and routine analyses of blood p-tau become routine practice in clinical assessments and research studies, probably combined with (1) high-performing assays of plasma A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> ratio for preclinical AD (Fig. 2), (2) brief digital cognitive testing for prognosis (Fig. 2) and (3) plasma NfL when suspecting non-AD neurodegenerative diseases. However, several outstanding challenges must be addressed. We need to obtain analytical standardization and quality control to provide a framework with which biotechnical companies and clinical laboratories can ascertain that they produce valid biomarker results. We need to demonstrate biomarker validity in diverse cohorts. Finally, we need to perform studies to prospectively generate real-world clinical data on the performance of blood-based AD biomarkers, especially in primary care settings. We do not yet know how observations from such cohorts will translate to the setting of routine memory clinics, which see patients with greater heterogeneity in demographics, disease presentation and comorbidities. Therefore, whether blood p-tau can be used as a single marker or replace CSF biomarkers that have been tested in larger varieties of disease conditions remains unclear. Realistically, we might need to exercise caution in projecting immediate diagnostic use of blood p-tau levels as a CSF substitute until large-scale clinical characterization studies have been performed. Finally, we want to stress that we need to develop blood biomarkers for non-AD brain pathologies, especially for pathological changes in TDP-43, 3R tau, 4R tau,  $\alpha$ -synuclein and cerebrovascular changes as well as synaptic dysfunction.

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**Box 1**

**Factors governing the clinical robustness of a biomarker**

Total measurement variability		Clinical biomarker performance		
Factors that may lead to variability in measured concentrations include		The clinical diagnostic performance of a biomarker depends on		
<b>Biological factors</b>	<b>Patient-centered factors</b>	Differences across patients may affect biomarker levels. Examples: age, sex, race/ethnicity, genetics (for example, <i>APOE</i> genotype), obesity, comorbidities (for example, kidney dysfunction), medication (for example, enzyme inhibitors).	<b>Biomarker effect size</b>	The effect size, either an increase or a decrease, of the biomarker is assessed using, for example Cohen's <i>d</i> , defined as the difference in mean between patients with and without (unaffected control individuals or patients with other diseases) the disease or pathology (for example, brain amyloidosis), divided by the pooled standard deviation.
	<b>Within-individual factors</b>	Temporary influences on biomarker levels in an individual may include hydration status, diurnal variability, stress and concurrent minor infections.		The effect size is linked to other clinical biomarker performance characteristics including Sensitivity (the probability that the test is positive in those who have the disease or pathology) Specificity (the probability that the test is negative in those without the disease or pathology) Positive predictive value (the probability that a patient has the disease or pathology when the test is positive) Negative predictive value (the probability that a patient does not have the disease or pathology when the test is negative)
<b>Technical and analytical factors</b>	<b>Pre-analytical factors</b>	Fluid-collection and -processing methods may affect the measured level of the biomarker. Examples include type of collection tube, time to centrifugation, shipment time and temperature, and possible freeze-thaw.		The effect size for a biomarker may be smaller in patients with preclinical or early stages of the disease (with limited amounts of pathology) than in patients with



Total measurement variability		Clinical biomarker performance
Factors that may lead to variability in measured concentrations include		The clinical diagnostic performance of a biomarker depends on
	Pre-analytical conditions may vary between samples, between centers and over the season.	advanced disease (and extensive pathology).
<b>Analytical performance of biomarker assay</b>	Analytical variability means the inevitable differences in concentrations measured present for any measurement technique, both within a run and between runs. Analytical variability is measured through fit-for-purpose validation experiments and includes, for example, within-run precision, between-run repeatability and accuracy (which can be assessed when reference standards exist).	
<b>Bias</b>	Differences or drifts in measured concentrations between laboratories, batches of reagents or changes in analytical equipment (for example, liquid chromatography columns or instruments)	
Biomarker performance and robustness		
<b>Prospective biomarker studies</b>	Clinical biomarker performance characteristics depend on the conditions under which they were assessed. Biomarker performance will be higher in selected research cohorts (with a high proportion of typical patients with AD and healthy controls) than in unselected primary care or memory clinic cohorts and also depending on how samples were analyzed (batch analysis with identification of the optimal cutoff with the biomarker data in hand versus clinical routine-like multiple analyses with a pre-set cutoff). Thus, clinical biomarker performance characteristics need to be assessed in prospective clinical trials, in settings resembling the conditions in future clinical routine, meaning a clinical cohort representative of the intended use population and with analyses	

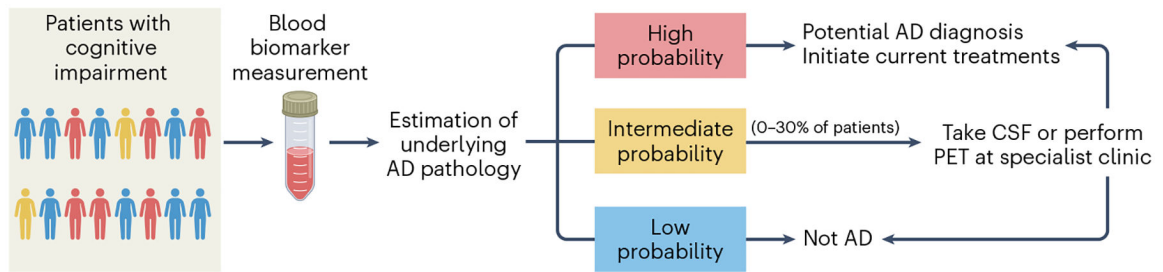
Total measurement variability		Clinical biomarker performance
Factors that may lead to variability in measured concentrations include		The clinical diagnostic performance of a biomarker depends on
	performed continuously and patients classified using a pre-set cutoff into those with the disease or pathology and those without.	
<b>Clinically robust biomarker</b>	A clinically robust biomarker has a total measurement error that is substantially lower than the biomarker effect size. This biomarker gives an accurate and consistent classification of patients into those who have the disease (or pathology, for example, brain amyloidosis) and those who do not.	

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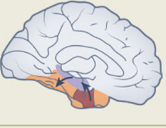


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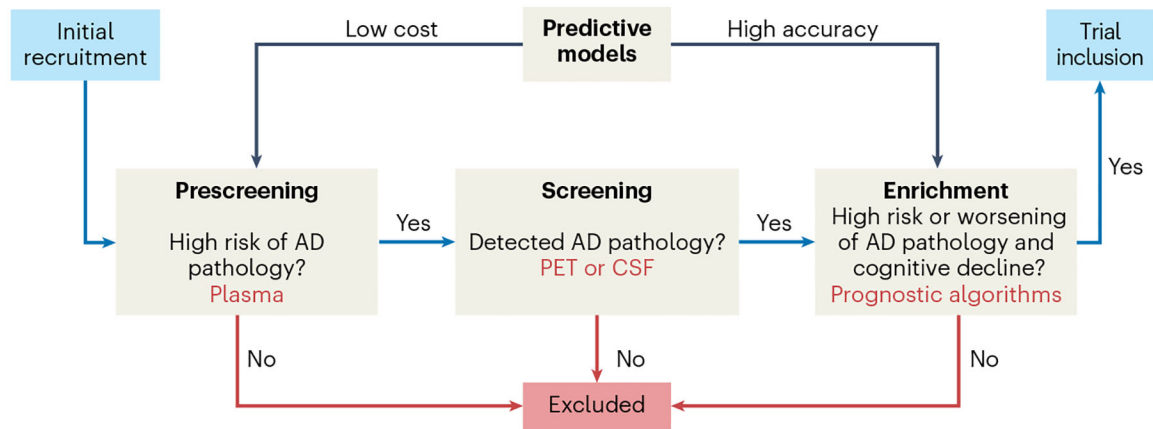


**Fig. 1 | Suggested blood-based biomarker-based workflow for Alzheimer's disease diagnostics.** Patients with cognitive complaints undergo blood sampling as part of the standard diagnostic work-up. High-performing blood Alzheimer's disease (AD) biomarkers (for example, p-tau217) are used to determine the individual-level probability of having AD. For patients deemed to have a very low probability based on blood-based biomarkers (BBMs), another cause of the symptomatology should be sought. For patients deemed to have a very high probability based on BBMs, appropriate treatments might be initiated. Patients with an intermediate probability, whose BBM results lie in an uncertain 'gray zone', might be referred for confirmatory testing with either cerebrospinal fluid (CSF) or positron emission tomography (PET) AD biomarkers. The percentage of individuals in such a 'gray zone' will depend on the accuracy of the blood-based diagnostic algorithm (very-high-performing BBM assays will have few results ending up in the 'gray zone').

	Cognitively unimpaired	MCI	Dementia
A $\beta$ pathology			
Tau pathology			
AD diagnosis	p-tau231 (or p-tau217) + A $\beta$ <sub>42</sub> /A $\beta$ <sub>40</sub>	p-tau217 (+MTBR-tau)	p-tau217 (+MTBR-tau)
Predicting AD dementia	p-tau217 + cognitive tests (+MTBR-tau)	p-tau217 + cognitive tests (+MTBR-tau)	NA

**Fig. 2 | An overview of key blood-based biomarkers used in the diagnostic or prognostic work-up of Alzheimer's disease.**

The top two rows depict the evolution of amyloid- $\beta$  (A $\beta$ ) and tau pathological brain changes during the different disease stages of Alzheimer's disease (AD)<sup>2</sup>. The third row shows that high-performing p-tau217 assays will probably be sufficient for detection of AD brain pathological changes in patients with cognitive impairment (mild cognitive impairment (MCI) or dementia)<sup>54</sup>. However, during the preclinical stages of the disease (when individuals are still cognitively unimpaired), p-tau231 and A $\beta$ <sub>42</sub>/A $\beta$ <sub>40</sub> are especially important to detect AD brain changes<sup>61</sup>. The bottom row shows that plasma p-tau217 is very important for high-performing prognostic algorithms predicting subsequent development of AD dementia in cognitively unimpaired individuals and patients with MCI<sup>90,98</sup>. Furthermore, in such prognostic algorithms, brief cognitive tests also contribute to the predictive accuracy. Future studies are needed to determine whether new fluid tau markers such as microtubule-binding region (MTBR)-tau will add diagnostic and prognostic information when combined with the already established markers. NA, not applicable.



**Fig. 3 |. Suggested workflow for inclusion of study participants into preclinical Alzheimer's disease trials.**

In the 'prescreening' step, a diagnostic algorithm based on blood-based biomarkers (BBMs) for Alzheimer's disease (AD) identifies cognitively normal individuals as being at low risk or high risk of having pre-symptomatic (preclinical) AD. In the 'screening' step, individuals deemed high risk will undergo further tests, involving amyloid- $\beta$  ( $A\beta$ ) positron emission tomography (PET) or cerebrospinal fluid (CSF) AD biomarkers, to confirm or rule out the presence of AD pathology. The prescreening step with BBMs will result in substantial time and cost savings, as far fewer CSF or PET tests will be needed to identify a certain number of individuals with preclinical AD. In the 'enrichment' step, a prognostic algorithm can be used to identify individuals who are likely to subsequently exhibit more severe spread of tau pathology and cognitive decline, so that the population to be included in the trial is enriched for such individuals. This latter enrichment step enables preclinical AD trials with shorter durations and/or fewer study participants.

**Table 1 |**

Current overview of candidate biomarkers with relevance to Alzheimer's disease

	<b>Fluid biomarkers</b>	<b>Imaging biomarkers</b>
<b>A<math>\beta</math></b>	Clinical: A $\beta$ <sub>42</sub> , A $\beta$ <sub>42</sub> /A $\beta$ <sub>40</sub> , A $\beta$ <sub>42</sub> /p-tau Experimental: oligomers of A $\beta$	Clinical: A $\beta$ PET imaging (for example, <sup>11</sup> C[PiB])
<b>AD-like tau</b>	Clinical: p-tau217, p-tau181 Experimental: p-tau231, p-tau212, p-tau205, MTBR-tau and others	Clinical: tau PET imaging (for example, <sup>18</sup> F[flortaucipir])
<b>Neurodegeneration</b>	Clinical: NfL Experimental: neurogranin, NPTX2, SNAP-25, GAP-43, $\beta$ -synuclein, 14-3-3 and others	Clinical: vMRI, FDG PET Experimental: dMRI, ASL
<b>Astrocytic response</b>	Clinical: GFAP Experimental: YKL-40	Experimental: deprenyl PET and others
<b>Microglial response</b>	Experimental: sTREM2, TAM receptors and others	Experimental: TSPO PET and others

The table divides biomarkers into those that can be measured in fluids (cerebrospinal fluid (CSF) and/or blood) and by brain imaging (positron emission tomography (PET) or magnetic resonance imaging (MRI)). It is important to note that not all fluid biomarkers are relevant to Alzheimer's disease (AD) when measured in blood but only when measured in CSF (such as neurogranin, soluble triggering receptor expressed on myeloid cells 2 (sTREM2) and YKL-40 (also known as chitinase-3-like protein 1 (CHI3L1)), because they are expressed to a high degree outside the brain as well. The table also indicates biomarkers that might be used in clinical practice and those that are still more experimental. A $\beta$ , amyloid- $\beta$ ; ASL, arterial spin labeling; dMRI, diffusion MRI; FDG PET, fluorodeoxyglucose PET; GAP-43, growth-associated protein, 43 kDa; GFAP, glial fibrillary acidic protein; MTBR, microtubule-binding region; NfL, neurofilament light; NPTX2, neuronal pentraxin 2; PiB, Pittsburgh compound B; SNAP-25, synaptosomal-associated protein, 25 kDa; TSPO, translocator protein, 18 kDa; vMRI, volumetric MRI.