

Neuroimaging markers for the prediction and early diagnosis of Alzheimer's disease dementia

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Alzheimer's disease (AD) is a progressive age-related neurodegenerative disease. At the time of clinical manifestation of dementia, significant irreversible brain damage is already present, rendering the diagnosis of AD at early stages of the disease an urgent prerequisite for therapeutic treatment to halt, or at least slow, disease progression. In this review, we discuss various neuroimaging measures that are proving to have potential value as biomarkers of AD pathology for the detection and prediction of AD before the onset of dementia. Recent studies that have identified AD-like structural and functional brain changes in elderly people who are cognitively within the normal range or who have mild cognitive impairment (MCI) are discussed. A dynamic sequence model of changes that occur in neuroimaging markers during the different disease stages is presented and the predictive value of multimodal neuroimaging for AD dementia is considered.

Introduction

An estimated 35.6 million people are affected worldwide today by a mild to severe clinical dementia syndrome, associated with costs of approximately US\$ 604 billion (World Alzheimer Report 2010, <http://www.alz.org>). The number of affected people is predicted to dramatically increase to 115 million by 2050 (World Alzheimer Report 2010, <http://www.alz.org>). Dementia of the Alzheimer's disease (AD) (see Glossary) type is the most frequent form of age-related dementia [1,2].

Clinically, initial progressive memory deficits that are eventually accompanied by more global cognitive and attention deficits are typical in AD dementia. Major pathologies in the brain associated with AD dementia have been identified, but the causes of such pathological changes are largely unknown. Causative factors include autosomal dominant inheritable mutations in the genes of presenilin

Glossary

Amyloid-beta (A β): the peptide that is the main component of amyloid deposits in forms of plaques in the brain, such as those detected by amyloid PET tracers. The A β peptide is cleaved at different lengths of amino acid chains (isoforms) from the amyloid precursor protein (APP), and the isoform A β ₁₋₄₂ shows increased tendency to aggregate into plaques. Although plaques are abundant in the AD brain, the exact mechanism of neurotoxicity of A β has not been established. Recent evidence suggests a role of the formation of soluble A β oligomers in neurodegeneration [71].

Alzheimer's disease (AD) dementia: according to the traditional NINCDS-ADRDA criteria, AD dementia includes gradual onset of cognitive impairment in episodic memory and at least one other domain [14]. The recent NIA-AA revision of the diagnostic criteria does not make impairment in the episodic memory domain mandatory anymore, but requires impairment in amnesic or nonamnesic cognitive abilities [15]. Presence of impaired social functioning and instrumental activities of daily living is considered as the crucial difference for the clinical diagnosis of AD, as opposed to MCI.

AD according to the Dubois criteria [11,12]: presence of episodic memory impairment and abnormality on at least one of the core MRI-, PET- or CSF-based biomarkers of AD pathology are required. Encompasses both subjects with abnormal biomarkers who show clinical AD dementia and those who only show mild memory impairment (i.e. prodromal AD).

Apolipoprotein E (ApoE) gene: the gene encodes, through three different alleles (ϵ 2, ϵ 3 and ϵ 4), three major isoforms of the ApoE protein, which are known to be involved in lipid metabolism. The ApoE ϵ 3/ ϵ 3 genotype is the most common form (prevalent in ~50–70% of the population). The ApoE ϵ 4 allele, which only occurs in approximately 10–15% of the population, accounts for the majority of AD cases [108] and is also associated with increased risk of other forms of dementia, such as vascular dementia and Lewy body dementia [109]. ApoE ϵ 4 shows a gene dose effect in that presence of two ApoE ϵ 4 alleles is associated with a higher risk compared with a single ApoE ϵ 4 allele [109]. In AD, the ApoE ϵ 4 isoform has been associated with increased A β deposition, although A β -independent mechanisms of ApoE, such as reduced adult neurogenesis in the hippocampus or phosphorylation of the protein tau (associated with increased tau pathology in AD), have also been demonstrated [108].

Biomarker for AD: a quantifiable surrogate measure of brain pathologies underlying AD dementia [110]. Neuroimaging-derived measures have been proposed as core biomarkers for the early detection of AD. Other types of biomarker candidates that are considered as core biological indicators of AD pathology, include CSF-derived measures of A β ₁₋₄₂, or the protein tau as a measure of neurofibrillary tangles (reviewed in [111]).

Clinical dementia rating (CDR): clinical rating scale used by the physician to assess performance in five different cognitive domains based on a semi-structured interview of both the patient and the informant [112]. The global CDR score ranges from 1 to 5, where a score of 0 means no cognitive impairment and a score of 5 corresponds to severe dementia. A score of 0.5 is often used in the diagnosis of MCI in clinical research studies [113].

Default mode network: during resting state, i.e. when a person is not challenged by external stimulation (e.g. a memory task), a brain circuit called the default

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mode network including the medial frontal, temporal and parietal brain regions are activated in healthy subjects. This network typically demonstrates beneficial deactivations during memory encoding in healthy subjects. Alterations in the activation pattern of the default network are also observed in numerous neurological disorders, including in AD and schizophrenia, as well as in aged subjects [42].

Mild cognitive impairment (MCI) (core clinical criteria): clinical diagnosis that includes progressive cognitive decline in memory (amnesic MCI) or other cognitive domains (non-amnesic MCI). MCI is characterized by subjective cognitive impairment reported by subjects themselves or their relatives in addition to cognitive impairment as assessed by neuropsychological tests, without affecting social functioning and instrumental activities of daily living [114]. According to the core clinical criteria of MCI proposed by the NIA-AA work group [13], other potential causes of cognitive decline, including vascular, traumatic or medical causes, or presence of symptoms of neurodegenerative diseases other than AD should be sought to be excluded by the clinician [13].

MCI due to AD (research criteria): subjects in this group fulfill the core clinical criteria of MCI, but also show pathophysiology of AD as confirmed by biomarkers [13]. This dual clinico-pathological definition of MCI due to AD is similar to prodromal AD (see below), with the difference that only progressive episodic memory impairment in addition to abnormal biomarkers are required to meet the criteria for prodromal AD.

Preclinical AD: a term that is used to refer to clinically normal subjects with evidence of AD pathology as detected by biomarkers. Subjects might show subjective memory impairment or very subtle cognitive decline, but would not demonstrate impairment of cognitive ability as detectable by standard neuropsychological tests. Such a definition also includes subjects who have genetic mutations that cause AD but have not already developed cognitive impairment [12]. A classification scheme of progressive accumulation of AD brain changes in preclinical AD has recently been proposed by the NIA-AA work group [16]. Preclinical AD is a research concept and not recommended for clinical routine diagnosis. It is currently unknown what proportion of patients with preclinical AD go on to develop AD dementia.

Prodromal AD: diagnostic classification for research purposes that includes the presence of episodic memory impairment (without affecting normal social functioning and instrumental activities of daily living) and, in addition, abnormalities as assessed with one or more biomarkers [12].

1 (*PS1*) and presenilin 2 (*PS2*) and the amyloid precursor protein (*APP*). Such an inheritable form of AD (familial AD) is associated with early onset of the disease, typically before the age of 65 years, but accounts for approximately only 1% of all AD cases [3]. For AD without the presence of such known genetic causes (sporadic AD), a late onset of dementia (age >65 years) is typical. The most important risk factor for sporadic late onset AD is age, with the annual incidence of clinically-diagnosed AD dementia being <1% of adults between 65 and 69 years old, but >8% in adults 85 years and older [4]. Presence of the apolipoprotein E (*ApoE*) $\epsilon 4$ allele is the strongest genetic risk factor of sporadic AD [3]. Presence of at least one *ApoE* $\epsilon 4$ allele advances the age of clinical onset of AD dementia significantly (from age 84 to 68 years) and increases the risk of AD dementia by a factor of four (for regularly updated meta-analyses, see <http://www.alzgene.org/>).

Core neuropathologies in AD include abnormalities such as the accumulation of the protein amyloid-beta ($A\beta$) and the development of neurofibrillary tangles, which have been associated with neuronal degeneration and clinical symptoms of dementia [5]. Such brain changes occur decades before the onset of dementia. The detection of $A\beta$ in the brain of living subjects has been made possible by recent developments in positron emission tomography (PET) using radiotracers such as ^{11}C -labeled Pittsburgh Compound-B (PiB), which label $A\beta$ deposits [6]. Such studies have confirmed that substantial levels of $A\beta$ deposits are present in subjects before the onset of dementia or even before any overt signs of cognitive impairment (as discussed later in the review). Other early brain changes

include a decline in synaptic function, as assessed by [^{18}F]fluorodeoxyglucose positron emission tomography (FDG-PET), gross neuronal loss causing atrophy, as measured by volumetric magnetic resonance imaging (MRI), and white matter changes within axonal projections, as detected by diffusion tensor imaging (DTI) (Box 1). These different types of changes might evolve sequentially and relate to the development of cognitive impairment within the clinical course of AD [7].

Mild cognitive impairment (MCI) is a well-defined clinical syndrome, which includes deficits in memory or other cognitive abilities [8]. Subjects with MCI have a higher risk to progress to AD dementia [9], but a substantial proportion of MCI subjects remain stable for years or revert to normal, indicating that clinical MCI symptoms can also stem from non-AD related etiologies. This notion is substantiated by the finding that only a subset of patients with MCI display measurable amounts of $A\beta$ on PiB-PET scans (for a review, see [10]). To increase the detection of MCI due to underlying AD pathology, the combination of cognitive deficits (as present at the MCI stage) with abnormal values for biomarkers indicative of AD pathology (e.g. $A\beta$ levels as measured by PiB-PET) has been proposed as research diagnostic criteria of predementia AD, called 'MCI due to AD' or 'prodromal AD' [11–13]. A revision of the clinical diagnostic criteria of AD by the National Institute of Neurological and Communicative Disorders and Stroke (NINCDS) and the Alzheimer's Disease and Related Disorders Association (ADRDA) [14] has recently been proposed [11,13,15]. For clinical use, the National Institute of Aging (NIA) and Alzheimer's Association (AA) work group has recommended purely clinical and neuropsychological criteria for the diagnosis of AD dementia and MCI [13,15]. For the use as research diagnostic criteria of AD

Box 1. Neuroimaging methods commonly used to detect *in vivo* brain changes associated with neurodegeneration and cognitive decline in human subjects

- **PET**, in combination with *in vivo* amyloid imaging agents that bind to fibrillar $A\beta$ deposits in the brain, has been a valuable technique for visualizing and quantifying the deposition of $A\beta$ throughout the brain in living subjects. Current ligands include the ^{11}C -labeled radiotracer PiB or ^{18}F -labeled tracers such as [^{18}F]florbetaben, [^{18}F]florbetapir (also called [^{18}F]AV-45) and [^{18}F]flutemetamol.
- **FDG-PET** uses [^{18}F]fluorodeoxyglucose to detect changes in glucose metabolism and blood flow in the brain. Unlike fMRI (see below), FDG-PET is mostly performed during resting state of the subjects owing to its relatively poor temporal resolution.
- **Structural MRI** detects tissue changes in the grey matter and white matter. This technique is especially sensitive for grey matter volume changes due to gross neuronal loss and atrophy.
- **DTI** is an MRI technique that is used to assess microstructural brain changes within the white matter fiber tracts of the brain. The most commonly used index is FA, which is determined by the degree of directionality (anisotropy) of the movement of the water molecules. A reduced FA value is reflective of axonal degradation and myelin damage in the brain.
- **In fMRI**, the blood-oxygen-level-dependent (BOLD) signal is used to measure blood flow and blood oxygenation, which is believed to correlate with changes in neuronal activity at a time scale of a few seconds. fMRI has been used to assess cognitive task-related changes in brain activity and basal brain activity during resting state.

and MCI, however, the dual clinico-pathological definition based on clinical testing and biomarker measurements has been adopted [13,15], and awaits further validation before use in clinical routine diagnostics. It should be mentioned that MCI due to AD is a diagnostic label in view of clinical and biomarker evidence, but this diagnosis does not mean that AD pathology has necessarily caused the clinical symptoms or that subjects with MCI due to AD necessarily progress to AD dementia. Rather, it is a research concept that has not yet been fully validated.

The question of the clinical fate of subjects with AD-like brain changes becomes even more urgent at an earlier time point, i.e. the preclinical phase of AD [16], when such brain changes are present without any overt cognitive deficits. The extent to which subjects with preclinical AD progress to dementia is currently not known. Of note, it has not been established that all subjects with AD-like brain changes inevitably progress to AD dementia, and many subjects with preclinical AD die of natural age-related causes before any signs of dementia are ever observed.

In the current article, we review recent imaging studies that have examined brain changes in both the preclinical and MCI phase, and evaluate the utility of neuroimaging markers for the prediction of clinical progression in these two distinct phases. The association between different imaging modalities is shown for each of the stages, and stage-specific changes in neuroimaging markers are discussed with respect to the predictive value for cognitive decline and the progression from MCI to AD dementia.

Multimodal neuroimaging changes in AD dementia

Before discussing neuroimaging changes of relevance for the preclinical and MCI due to AD stages, it is helpful to begin by briefly overviewing the typical neuroimaging findings that have been observed over the past several years in patients with clinically manifest AD dementia. Multiple studies have determined that the pattern of A β deposits detected in the brains of living subjects diagnosed with AD dementia using [11 C]PiB-PET closely matches the pattern predicted by the histochemical detection of A β in postmortem brain tissue from subjects with clinical AD [6,10] (Figure 1). Another amyloid PET imaging tracer that has been recently developed, the 18 F-labeled radioligand florbetapir (also known as [18 F]AV-45), has shown similar results with good correspondence between the distribution of amyloid PET and histochemically detected A β in the brain [17]. Clinical utility is expected, especially from these 18 F-labeled ligands, which possess a radiotracer half-life that is sufficiently long enough to be produced off-site and shipped to clinics. However, compared to [11 C]PiB-PET, such novel amyloid PET ligands are currently less well characterized in clinical populations of MCI and AD patients [10].

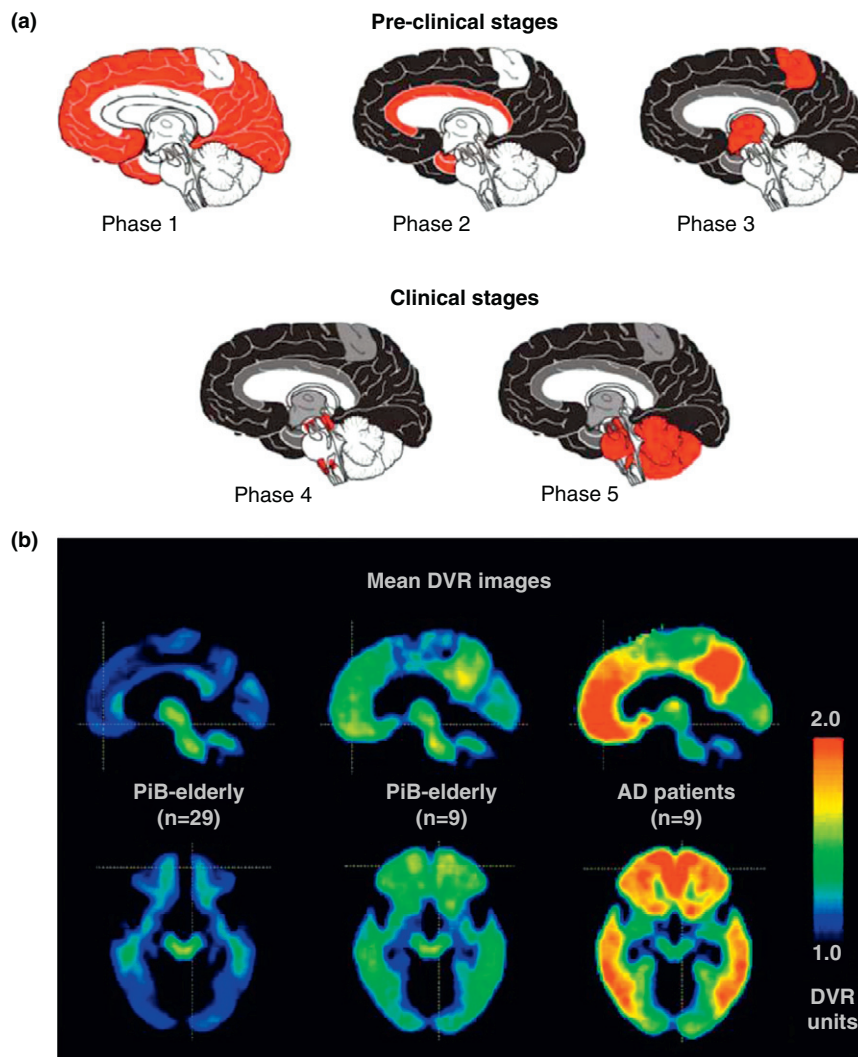
During the pre-clinical stages of AD, the pattern of A β in the brain, as derived from histochemical post-mortem brain examinations [18], shows an initial diffuse distribution within the neocortex and progresses towards the temporal allocortex, including the hippocampus, as well as subcortical brain structures during the course of the disease (Figure 1a). In the clinical stages of cognitive impairment, A β deposits are increased in both the amount

and spatial extent throughout the brain [18] (Figure 1a). At the stage of AD, A β deposits are most frequent in neocortical brain regions, hippocampus, amygdala and subcortical brain regions (e.g. striatum and basal ganglia), but only weakly distributed within the cerebellum and some brain stem areas [18]. Consistent with the histochemical post-mortem findings on A β deposition in the brain, a distribution of amyloid PET uptake can be already seen widely within the brain in the preclinical stage [19]. A significant increase in A β deposition in predilection brain areas of AD can be observed in AD dementia patients (Figure 1b). Based on global brain PiB-PET uptake, subjects are often dichotomized into groups with high and low PiB-PET values, i.e. PiB-PET(+) and PiB-PET(-) groups [19]. Data from 15 research groups have shown that 96% of 341 clinically-diagnosed AD patients and 24% of 651 cognitively normal elderly controls were PiB-PET(+) [19–28]. This finding highlights the diagnostic sensitivity of PiB-PET imaging, but also demonstrates that A β pathology is not specific for the dementia phase of AD, similar to what has been known from brain autopsy studies [29].

Longitudinal studies including serial scans provide an assessment of individual trajectories of brain changes and give insight into variability of brain changes between different persons. A recent longitudinal study with serial PiB-PET scans has reported a mean annual increase of 4% in global PiB-PET uptake over a period of 2 years in elderly subjects with clinically-diagnosed AD [30]. However, such changes are relatively small and show substantial variability between AD patients [31]. Furthermore, some studies do not report a significant increase in global PiB-PET uptake with time, at least when assessed over a few years [32–34], suggesting that A β deposition might increase only slowly or has already reached a plateau in patients with AD dementia [33]. However, it should be noted that, to date, most longitudinal studies have included only small numbers of subjects (i.e. usually <20 subjects), and owing to the substantial variability between subjects, such studies might have lacked sufficient statistical power to detect a significant annual increase in PiB-PET [32,33,35].

FDG-PET-derived measures of brain glucose metabolism and cerebral blood flow are markers of synaptic dysfunction, typically obtained during resting state (Box 1). A characteristic pattern of hypometabolism in the temporo-parietal region of the cortex, which is involved in episodic memory function, is present at the AD dementia stage [36,37]. Joint assessment of FDG-PET and PiB-PET shows the expected inverse association between both modalities within the temporo-parietal region, although not frontal regions [38,6], suggesting that hypometabolism in core brain regions is associated with A β pathology.

A well-established finding using functional MRI (fMRI) (Box 1) to examine AD-associated differences in brain activation is decreased activation in the hippocampus during episodic memory tasks [39,40]. Such results are consistent with clinical findings of early episodic memory deficits in AD subjects [41]. Abnormal fMRI-assessed brain activation can also be observed in AD subjects without engaging subjects in a cognitive task. In healthy subjects, the default network of brain regions is active during resting periods (probably a reflection of cognitive processes



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Figure 1. The progression of amyloid deposits as assessed at different stages of AD in (a) post-mortem brain tissue using immunohistochemical techniques against A β and (b) *in vivo* amyloid PET scans. (A) Phases (1–5) of postmortem histological appearance of A β in clinical AD (based on [18,115]). Note that the stages refer to A β brain deposition and not necessarily to clinical severity. In phase 1, A β is largely restricted to neocortical brain areas. In phases 2 and 3, when clinical signs of cognitive decline have yet to appear, A β deposits are localized throughout the neocortex, all cortex, including brain areas such as the medial temporal lobe, and also begin to occur in the motor cortex and subcortical brain structures. In the clinical stages of AD (phases 4 and 5), A β deposits are observed globally in the brain including the brain stem and the cerebellum in the final stage of the disease (phase 5). Key: white, no A β deposits; red, novel A β deposition; grey, A β deposits that were missing in phase 1; black, A β deposits that were present already in phase 1. (b) A similar spatial pattern of progression of A β deposition is observed when using PiB-PET in cognitively normal elderly controls who appear to be A β -free (PiB–), cognitively normal elderly controls who appear to be in the early stages of A β deposition (PiB+) and in symptomatic AD patients [19]. Mean sagittal distribution volume ratio (DVR; cerebellum as reference) images are at the top and transaxial images at the bottom. Reproduced, with permission, from [18,115] (a) and [19] (b).

such as introspection) but becomes deactivated during cognitive processes that are focused on external stimulation, such as performance of a cognitive task [42]. The intrinsic functional connectivity (i.e. the coordinated co-activation of the default network's brain regions measured with resting state fMRI) is impaired in AD dementia subjects during the resting state as compared to cognitively healthy elderly subjects [43,44]. Furthermore, the default mode network of AD subjects do not show the beneficial deactivations that healthy subjects show when assessed during memory tasks [45]. The abnormal fMRI activation and connectivity within the default mode network brain regions overlaps spatially with the temporo-parietal regions of FDG-PET hypometabolism in AD subjects, suggesting that both modalities exhibit a converging pattern of functional brain impairment in AD.

Structural MRI is another imaging technique that has been used to assess brain changes in AD subjects, specifically with respect to grey matter volume changes. A recent meta-analysis that included 826 patients with AD and 1027 elderly cognitively normal subjects found that the medial temporal lobe showed the strongest changes in AD dementia [46], with a volume loss of 20% in the hippocampus already present at a mild stage of AD dementia [46]. Other brain areas including the lateral temporal lobe, parietal and prefrontal lobes were also found to be negatively affected, but changes in brain regions outside of the medial temporal lobe were smaller and more variable across studies [47].

An assessment of white matter fibers using the imaging technique of DTI has revealed that the fibers connecting the hippocampus and posterior cingulate gyrus are im-

paired in AD subjects to a significantly greater degree compared with control subjects [48]. This suggests that white matter damage might relate to grey matter atrophy within the temporo-parietal brain network in AD. A recent meta-analysis of DTI studies of AD subjects confirmed a large effect size of white matter damage in the posterior cingulum, but also within major white matter bundles connecting the prefrontal cortex with the medial temporal lobe or the parietal cortex [49], suggesting that white matter damage affects large-scale networks in AD.

Taking such diverse imaging modality findings together, it is clear that gross changes in the living brain of AD dementia subjects can be visualized and measured. A hypothetical model of sequential neuroimaging changes in AD has recently been proposed [7]. Box 2 provides an overview of disease-stage specific changes for different neuroimaging markers throughout the course of the disease. The neuroimaging changes in the early stages of disease preceding dementia are discussed in the following sections.

Multimodal neuroimaging in preclinical AD

In this section, we will provide an overview of neuroimaging studies that have assessed changes in subjects with preclinical AD, i.e. in elderly subjects who show normal cognitive abilities but have already AD-like brain abnor-

malities such as discussed above. Findings from amyloid PET studies will be discussed first and compared to findings on amyloid PET in AD dementia. Functional and structural brain changes (as assessed by FDG-PET and MRI methods) will be discussed subsequently, especially in relation to early A β pathology, taking into account the influence of *ApoE* genotype as a potential modulating factor of such brain changes at this early disease stage.

Amyloid PET

As discussed above, PET imaging studies using the PiB ligand have revealed that about 10–30% of elderly cognitively normal subjects are PiB(+) [10]. *ApoE* genotype is known to modulate the level of A β deposition in the brain [22], and homozygous *ApoE* ϵ 4 carriers show the highest proportion of PiB(+) [50]. A recent study that imaged amyloid in the brains of subjects within the age range of 18 and 50 years using [18 F]AV-45-labeled PET revealed an absence of A β deposits, regardless of *ApoE* genotype [17]. Together, these findings suggest that substantial elevations in A β occur only after 50 years of age [17]. An increase in A β levels, as assessed by PiB-PET, has been associated with more rapid memory and global cognitive decline [51]; whether cognitively normal subjects with increased PiB-PET uptake will show higher likelihood to progress to

Box 2. Biomarkers commonly used to detect the development of AD dementia

(1) Neuroimaging biomarkers

There are a variety of neuroimaging biomarkers in use for the prediction and validation of AD. Some of these techniques detect core AD pathology such as amyloid deposition (as measured by amyloid PET), whereas other imaging modalities detect neurodegeneration such as FDG-PET- and MRI-assessed functional and structural changes that might occur in a specific temporal order as recently proposed [7]. Some neuroimaging-detected changes occur already at the preclinical stage, others at the MCI stage and all of them are useful for detecting changes at the AD dementia stage (Figure 1). The dynamic changes in neuroimaging markers are known to occur non-linearly throughout the stages of the disease [116,117]. However, the exact temporal sequence of structural and functional brain changes, and how changes in these various different imaging modalities are

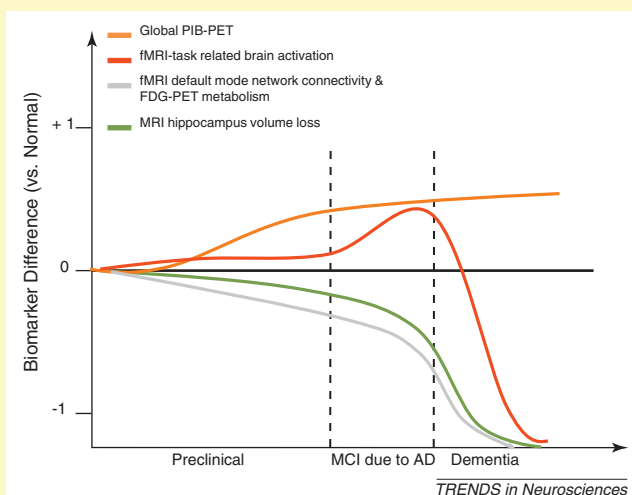


Figure 1. Hypothetical model of various neuroimaging biomarkers and their predicted utility during disease progression, as based on a variety of studies discussed in this review.

related, remains to be determined. The sequence model displayed presents a hypothetical framework based upon currently available neuroimaging data.

- (i) *PiB-PET imaging* (orange line): studies have observed that 10–30% of elderly cognitively normal subjects have significant A β deposition (i.e. already at the preclinical phase) [10].
- (ii) *fMRI task-related brain activation* (red line): increased PiB-PET is correlated with abnormally increased hippocampus activity during memory task performance. Such functional changes begin to be observed in the late preclinical stage and decline late in MCI [60,90]. Abnormally decreased hippocampal activity is observed in this later stage [41,45].
- (iii) *fMRI-assessed default mode network* (grey line): begins to decline in activation levels during the preclinical stage, in correlation with PiB-PET [23,60,62]. Resting state FDG-PET metabolism (grey line) is reduced in *ApoE* ϵ 4 carriers without cognitive impairment that might, however, not be accounted for by preclinical deposition of A β pathology [118].
- (iv) *Volumetric MRI* (green line): MRI-detected grey matter atrophy starts primarily, though not exclusively, in the hippocampus, in inverse correlation with PiB-PET levels, and continues to decline throughout the progression of the disease [68,117]. Widespread cortical and subcortical atrophy is observed at the dementia stage [119].

(2) Other biological-based biomarkers

Another major category of primary biomarker candidates for AD consists of CSF biomarkers (for review, see [111]). CSF samples can be obtained from subjects by lumbar puncture and analyzed in standard laboratory tests. Major CSF-derived markers include:

- (i) *Soluble A β _{1–42}*, which correlates inversely with A β plaque deposition in the brain [120] and global PiB-PET scores [121];
- (ii) *Phospho-tau*, which correlates positively with neurofibrillary pathology in the brain [122];
- (iii) *Total tau*, which is thought to be associated with neuronal loss in the brain because it is elevated in neurodegenerative diseases such as Creutzfeldt-Jakob disease or stroke, which show large neuronal loss but are inconspicuous of AD-like neurofibrillary pathology (for reviews, see [123,124]).

dementia is, however, unknown and studies to date have not included sufficient follow-up durations to adequately assess this question. Vulnerability factors (such as sub-clinical cerebrovascular disease) and brain/cognitive reserve factors are likely to play a modulatory role in the process of A β deposition [27,52–54]. However, the strongest determining factor at this stage remains the *ApoE4* allele, therefore this genetic information is important to take into account when examining functional and structural brain changes in subjects.

FDG-PET

A recent study of cognitively normal subjects who were characterized as PiB-PET(+) did not find any evidence of FDG-PET hypometabolism in the prefrontal cortex and parietal lobe during resting state [55]. However, additional studies are needed to replicate this finding and to assess whether such a conclusion is common across different experimental groups. Other studies that have assessed FDG-PET hypometabolism in a subset of cognitively normal subjects that carry the *ApoE* ϵ 4 allele have detected changes in a subset of regions affected in AD [56,57], and these changes predict progression to AD [58]. Since *ApoE* ϵ 4 already increases the deposition of A β in the preclinical stage [50], it is possible that such genetic effects on FDG-PET are related to the presence of A β pathology. However, results from FDG-PET data obtained in *ApoE* ϵ 4 carriers within the 20–39 year age range suggest that a fundamental metabolic abnormality is already present at an age when substantial fibrillar A β deposition is unlikely to have developed [59]. This suggests that *ApoE* genotype could exert an influence on FDG-PET metabolism beyond its influence on A β in the preclinical phase.

FMRI

FMRI studies have uncovered several AD-like alterations in brain activity that are associated with A β or *ApoE* ϵ 4 genotype at an early preclinical phase. Although memory task-induced brain activation within the medial temporal memory system does not appear to be strongly affected in cognitively normal PiB-PET(+) subjects, the default network shows an AD-characteristic lack of deactivation, and even reverts into paradoxical increases in some of the network's brain areas [60,61]. In further parallel to the findings in AD dementia, the functional connectivity between brain regions of the default mode network is disrupted in elderly PiB-PET(+) adults [23,62,63]. These findings suggest that A β is associated with abnormal regulation of large-scale networks, especially within the default mode brain regions, even at this early preclinical stage [60]. However, disruption of default mode network connectivity has recently been reported in elderly adults who carry the *ApoE* ϵ 4 allele but who are PiB-PET(–) [64]. Furthermore, the presence of the *ApoE* ϵ 4 allele has also been associated with increased cortical activation during a visual learning task [65] and disruption of functional connectivity of the default mode network [66]. Therefore, similar to the findings of an *ApoE* effect on FDG-PET at the preclinical stage, *ApoE* ϵ 4 genotype and elevation of A β might both influence, to some extent independently, brain activation [64].

Structural MRI

There is now accumulating evidence that A β deposition in the brain is associated with grey matter atrophy in the preclinical stage of AD. In the largest combined cross-sectional MRI and PiB-PET study to date that included 135 elderly cognitively normal subjects, PiB(+) subjects showed higher hippocampus and cingulate cortex atrophy compared with PiB(–) subjects [67]. Furthermore, clinical longitudinal assessment revealed that PiB(+) subjects, but not PiB(–) subjects, showed decline in episodic memory and working memory over a time span of 16 years that preceded or succeeded the PET scan [67]. Consistent with these findings, other cross-sectional studies have shown an association between PiB-PET and hippocampus volume loss [68,69] and posterior cingulate gyrus atrophy [70]. A major question is whether there is a threshold level of A β deposition at which brain atrophy occurs. Some researchers argue that even small levels of elevation in A β , including levels that are currently defined as PiB(–), could already be associated with increased brain atrophy [68]. By contrast, the observation in PiB-PET studies that cognitively normal subjects show a bimodal segregation into groups of high and low levels of PiB binding [i.e. PiB(+) and PiB(–)] [35] suggests the existence of a distinct group of elderly adults with abnormal and potentially disease-specific elevations of A β . At this stage, the exact definition of pathological levels of A β deposition remains open. Furthermore, other pathological events such as the formation of soluble A β oligomers [71] or the development of tau-related neuropathologies [72] are likely to be critical factors in A β -related neurodegeneration, and will need to be taken into consideration in future assessments of A β -associated brain atrophy.

ApoE ϵ 4 genotype could be an important determinant also for brain atrophy at the preclinical stage. Similar to cross-sectional findings in AD dementia subjects [73], elderly cognitively normal subjects who carry the *ApoE* ϵ 4 allele have greater hippocampus volume loss when compared to elderly cognitive normal adults without that allele [74]. A large longitudinal population-based study with serial MRI scans in 1186 healthy subjects showed that cognitively normal subjects with two *ApoE* ϵ 4 alleles (i.e. homozygous) had a faster atrophy rate for the hippocampus and whole brain over a 4-year interval compared with heterozygous *ApoE* ϵ 4 carriers and non-carriers [75]. Therefore, both elevated A β levels and *ApoE* genetic risk are already associated with grey matter atrophy in subjects without cognitive impairment.

To date, microstructural white matter changes, such as those detected by DTI, have not been assessed in preclinical subjects in association with PiB-PET, and only few studies have investigated the impact of *ApoE* genotype on white matter changes in this subject group. Cognitively normal adults with at least one copy of *ApoE* ϵ 4 show reduced white matter integrity, as measured by a reduced fractional anisotropy (FA) value (Box 1) in the posterior cingulum, corpus callosum and other major white matter bundles that connect different lobes of the brain and support the default mode network [74]. These changes were observed in young adults (20–35 years) [76], middle aged (49–65 years) [77] and elderly cognitively normal

subjects (>65 years) [76]. Such changes are reminiscent of the early functional brain differences that were observed in ApoE $\epsilon 4$ carriers at both a young and old age, suggesting that these changes might occur independently of A β , as discussed above.

In summary, the amyloid PET findings are in accordance with histochemical autopsy studies demonstrating substantial deposition in the neocortex in elderly subjects without cognitive deficits, who are in the preclinical stages of AD. ApoE $\epsilon 4$ genotype contributes to A β deposition and influences brain function and brain anatomy. PiB-PET(+) subjects already display a number of structural and functional brain alterations including reduced functional connectivity between brain regions of the default mode network and accelerated medial temporal lobe atrophy that is associated with episodic memory impairment. Such functional and structural brain changes in PiB-PET(+) subjects are reminiscent of brain changes seen in AD dementia. Whether these early brain changes at the preclinical stage herald clinical progression to AD dementia is, however, currently unknown. In the next section, we review neuroimaging findings in subjects who are at a clinically more progressed stage, i.e. MCI.

Multimodal neuroimaging changes in MCI

In the following section, we review neuroimaging findings in subjects with clinically manifest MCI without full-blown dementia. The same imaging modalities as discussed previously are reviewed to allow for a direct comparison with findings in the preclinical AD and AD dementia stages discussed above.

Amyloid PET

Approximately 33–61% of subjects diagnosed with MCI show AD pathophysiology by virtue of being PiB-PET(+) [78]. The proportion of PiB-PET(+) subjects is intermediate compared to the proportion of PiB-PET(+) subjects found at the preclinical stage and the AD dementia stage (see above), suggesting an increase in the frequency of PiB-PET(+) from preclinical to clinical AD dementia stages. Clinical longitudinal studies show that subjects with MCI who convert to AD dementia within 2–3 years have higher baseline PiB-PET levels than MCI subjects who do not clinically worsen within that time frame [78,79]. The evidence for the utility of PiB-PET in identifying AD pathophysiology in the setting of clinical MCI is becoming increasingly convincing. When pooling data from nine PiB-PET studies that examined the PIB-PET status in MCI [21,24,28,79,80], overall, 161 of 272 MCI subjects were amyloid-positive (i.e. 59%). Five of these studies (155 MCI patients) included longitudinal clinical follow-up assessments, which showed that between 20 and 50% of subjects progressed to clinical AD over 1–3 years; the large majority of these MCI–AD converters were PiB-PET(+) at baseline [21,24,79–81]. This number will almost certainly increase as the follow-up time is increased to at least 5 years for all subjects. By contrast, only a relatively small proportion of PiB-PET(–) MCI patients (~8–15%) progressed to clinical AD in these studies [21,24,79–81]. Among PiB-PET(+) subjects with MCI, it is known that the ApoE $\epsilon 4$ genotype accelerates the time to progression

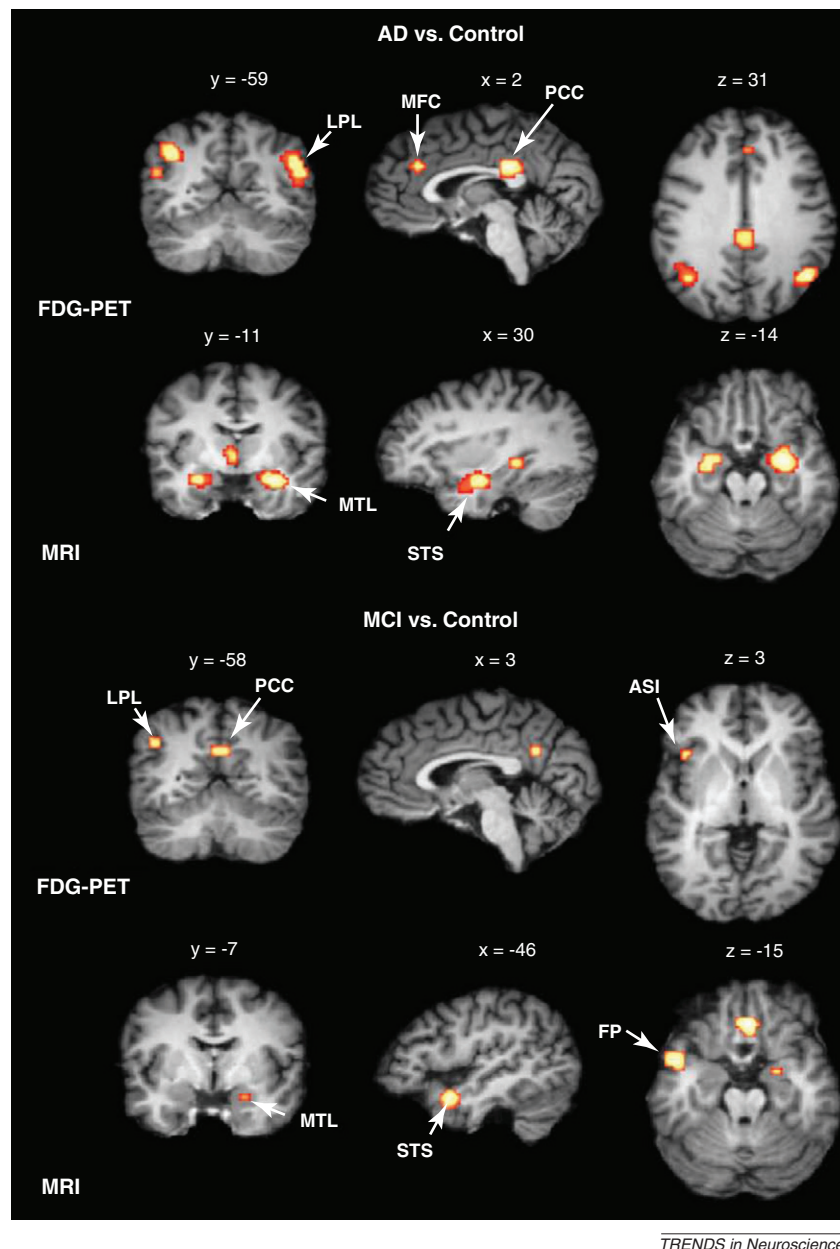
from MCI to AD [79]. Therefore, the combination of amyloid imaging together with genetic determination of ApoE alleles is valuable for predicting higher risk of progression to AD at the MCI stage.

FDG-PET

The AD-like hypometabolic pattern (e.g. decreased signals in the precuneus and temporo-parietal cortex) has likewise been observed in MCI subjects (for a review, see [82]; Figure 2). Indeed, several studies have shown that abnormalities in FDG-PET predict progression from MCI to AD [83–85]. Based on FDG-PET metabolism within the temporo-parietal brain regions and posterior cingulate gyrus, the test accuracy to predict conversion from MCI to AD dementia within 1–1.5 years ranged between 80 and 90% among these three studies, and both the sensitivity and specificity to predict AD dementia were above 80% in each study [83–85]. The predictive accuracy was even higher in ApoE $\epsilon 4$ carriers [85], suggesting that ApoE genotype might provide additional clinically relevant information for the prediction of AD dementia by FDG-PET hypometabolism. Whether or not FDG-PET hypometabolism in MCI is linked to deposition of A β is not clear at this stage. Only a few studies have examined the association between FDG-PET and amyloid PET at the MCI stage, reporting weak negative associations between FDG-PET and PIB-PET [86], which is consistent with the observation that FDG-PET and PIB-PET provide complimentary information for the prediction of conversion from MCI to AD [87].

fMRI

During resting state, disrupted connectivity of the default mode network is present in MCI [88], similar to what was reported in the preclinical stage in PiB-PET(+) subjects and subjects with AD dementia (as discussed above). Reduced default mode network connectivity has been associated with a higher risk of converting to AD dementia within a 3–4 year follow-up interval [89], suggesting that impaired default mode activity is predictive of impending clinical progression to dementia. Cognitive task-associated changes in fMRI have also shown to be predictive of cognitive decline in MCI [90]. Specifically, hyperactivation within the medial temporal lobe has been observed in the mild stages of MCI [45,91] in association with subsequent cognitive decline [90]. Such hippocampus hyperactivation during memory performance is already present in PIB-PET(+) subjects with a clinical dementia rating (CDR) of 0.5, who are in the transition phase between preclinical AD and MCI due to AD [60]. This suggests that memory-related hyperactivation of specific brain regions becomes abnormal early in the course of the disease. Interestingly, at clinically more advanced stages of MCI and in AD dementia subjects, this memory task-induced hyperactivation of the temporal-parietal memory network, however, turns into an activation deficit [45]. Specifically what this initial hyperactivation is due to and how that turns into an activation deficit remains unknown at this time; however, it could represent a transient phase of impending breakdown of neuronal networks. Possibly, the hyperactivation of particular brain areas reflects mechanisms of compensation within neural circuits. Alternatively, it could be



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Figure 2. Brain regions of functional and structural brain abnormalities that show strongest effects for the diagnosis and prognosis of AD dementia. Yellow-to-red labeled clusters show significantly decreased activity, as measured by FDG-PET (first and third row), and reduced grey matter volume, as measured by structural MRI (second and last row). AD subjects (top half) and MCI subjects (bottom half) were compared to elderly cognitively normal control subjects in a meta-analysis of voxel-based MRI and FDG-PET studies [47]. Note that in AD subjects, reduced FDG-PET is present in clusters of the posterior cingulum cortex (PCC), lateral parietal lobe (LPL) and medial frontal cortex (MFC), and is already present in the inferior PLP and PCC at the stage of MCI, in addition to a small cluster in the anterior superior insula (ASI). The majority of the grey matter volume decreases in AD appear in the medial temporal lobe (MTL) extending towards the superior temporal sulcus (STS), and these volume reductions are already present albeit to a lesser extent in MCI subjects. Clusters of significant group differences were projected onto standardized MRI images of brain slices in coronal view (left column), sagittal view (middle column) and axial view (right column). The x, y and z values label the slice coordinates within the standard Talairach atlas space. Adapted, with permission, from [47].

more directly related to A β pathology, either as a consequence of A β -induced hyperexcitability of neurons [92,93] or by increasing vulnerability to A β toxicity [94]. Whatever the underlying reason, initial evidence suggests that hyperactivity might reflect impending neuronal dysfunction and clinical decline [90].

Structural MRI

Several studies have shown that increased brain levels of A β (as assessed by PiB-PET), or decreased cerebrospinal fluid (CSF) A β levels, are associated with atrophy of the

hippocampus and other brain structures [68,95] in subjects with MCI, as assessed by volumetric MRI. These findings are consistent with those observed in preclinical AD. Multivariate statistical analyses have shown that an AD-typical pattern of MRI-assessed brain atrophy is present in a subgroup of MCI subjects [96,97]. Specifically, an atrophy pattern including the medial temporal lobe, posterior cingulate and orbitofrontal cortex is predictive of the development of AD dementia in MCI [97].

Apart from grey matter volume changes, increased white matter abnormalities have been observed in MCI.

DTI abnormalities within the medial temporo-parietal network associated with episodic memory impairment have been repeatedly reported in the literature (for a review, see [98]) and are among the brain regions that show the strongest predictive value for the discrimination between MCI and cognitively normal subjects [49,99,100]. Currently, there are only a few clinical longitudinal studies available for DTI [101–103], however, a recent study has shown that DTI-assessed diffusivity of the left hippocampus is predictive of the conversion from MCI to AD within 1.5 years [104].

In summary, AD-typical structural and functional changes at the MCI stage are reflected by changes in a variety of measures. Such measures include significant levels of amyloid PET binding in the brain, fMRI-assessed task-related medial temporal lobe hyperactivation and default network dysfunction, FDG-PET-assessed temporo-parietal hypometabolism, MRI-assessed medial temporal lobe atrophy and diffusion changes. In the next section, we address the question whether the combination of such markers is beneficial in the prediction of the risk of developing AD dementia within a few years.

Combining neuroimaging markers for the prediction of clinical progression from MCI to AD dementia

There is now accumulating evidence that a combination of neuroimaging markers show additive effects of different modalities for the prediction of progression to AD dementia. Cross-sectional studies suggest that PiB-PET and hippocampus volume provide complementary information for the diagnostic classification of AD dementia [35]. Within MCI subjects, the combination of structural MRI to assess hippocampus volume and DTI in the posterior parietal lobe contributes to the prediction of the severity of the memory deficits [102]. Neuroimaging studies with longitudinal clinical follow-up have shown that the combination of MRI-assessed grey matter atrophy and FDG-PET hypometabolism within the posterior cingulate is associated with increased risk of clinical progression within 2 years when compared with the use of either predictor alone [105]. When neuropsychological measures are also included as predictors, MCI subjects who are abnormal on both FDG-PET and episodic memory are approximately 12-fold more likely to progress to AD than subjects who were normal on these measures, which is significantly higher when compared to an increase in risk by approximately fourfold based on episodic memory deficits alone [106]. In that study, the hippocampus volume did not add to the predictive accuracy, suggesting redundancies between the different predictors tested. In fact, the prediction accuracy of some of the best single predictors including right entorhinal cortical thickness, right hippocampus volume or trail making test B (TMT-B) might only be marginally improved by combining multiple markers [107]. Therefore, although the risk prediction of AD might be increased by combining markers, case-by-case decisions of classifying subjects into MCI to AD converters might not necessarily be improved by the use of multiple neuroimaging markers. For the assessment of the utility of neuropsychological predictors as discussed above, it should be taken into account that neuropsychological performance is part of the definition of

Box 3. Outstanding questions

- Can early functional outcome measures or surrogate biomarkers (e.g. based on fMRI) be developed for use at early stages of disease progression, i.e. stages prior to established neurodegeneration? If so, is it possible to fully reverse such early changes, or at least halt disease progression, using disease-modifying treatments?
- What are reliable predictors of progression to AD dementia in cognitively normal subjects with high levels of A β in the brain?
- Are fMRI hyperactivation within the medial temporal lobe and disrupted functional connectivity within the default mode network reliable predictors of AD dementia and what is the underlying pathophysiological mechanism?
- What are reliable threshold values (i.e. quantitative decision criteria) for each of the different neuroimaging biomarker methods that can be used by clinicians to estimate the risk of AD in non-demented subjects and to diagnose AD dementia?
- What is the most cost-effective multi-biomarker model for the prediction of AD dementia?
- Is there a distinctive multi-modal neuroimaging marker signature from the first adaptational and functional brain changes to fully established neurodegeneration and dementia in AD subjects?

AD dementia, and thus neuropsychological predictors might correlate with progression to dementia, especially in studies that use only short clinical follow-up time intervals.

Eventually, any gain in predictive accuracy of a biomarker or test for the detection of AD needs to be weighed against costs in terms of side effects (e.g. radioactivity of PET tracers), potential invasiveness of methods (e.g. lumbar puncture in the case of CSF-based tests; Box 2) and availability of the technology (e.g. the imaging scanners and/or radiotracers might only be available at major hospitals and research centers). Notably, there are also psychological costs for the individual consequent to learning about the presence of AD-like brain pathology and possible clinical prognosis based on such biomarker test results. Such considerations are especially important to carefully weigh in light of the fact that no cure for AD or treatment to effectively slow or halt disease progression currently exists. At this stage, the main utility of biomarker-based early detection of AD is in identifying subjects with high likelihood of progression to AD dementia in clinical trials for the testing of novel drugs for disease prevention or treatment. Large-scale cost effectiveness studies are needed to evaluate the utility of biomarker-aided prediction of AD in clinical practice, and additional studies to better understand the underlying neurobiological changes that are occurring during disease progression in AD are needed (Box 3).

Concluding remarks

Neuroimaging methods are capable of detecting substantial brain changes, not only in subjects with AD dementia, but also in subjects in the mildly symptomatic MCI due to AD stage and even in cognitively normal subjects who might be in the preclinical stage of AD (Box 2). There are imaging modality-specific changes within these clinical/preclinical stages. In the preclinical stage, A β deposits are already present in a substantial number of subjects. Such changes are associated with increased grey matter brain atrophy, especially within the hippocampus, a key region affected in AD dementia. FMRI-assessed resting state functional connectivity and reduced deactivation of

the default network is already impaired in association with A β deposition in the preclinical phase. Hyperactivation within the hippocampus memory network during memory performance occurs early in the MCI phase but reduced hippocampus activation is visible only shortly before progressing to dementia, suggesting that such hippocampus hyperactivation is a transient sign impending clinical worsening. In contrast to fMRI-detected changes in brain activity, FDG-PET abnormalities have not yet been associated with A β deposition in the preclinical phase and only become detectable in the MCI phase. Therefore, resting state fMRI might be a more sensitive measure than FDG-PET to detect early changes associated with AD pathology. To date, DTI lacks a clear characterization at the preclinical stage but, similar to the other neuroimaging markers, shows specific patterns of alteration in association with genetic risk of AD. Across the various imaging modalities, *ApoE* genotype is an important modulating factor. The presence of the *ApoE* $\epsilon 4$ allele is associated with accelerating A β deposition, hippocampus atrophy and default mode network dysfunction, beginning in the preclinical stage.

At the stage of MCI, AD-like patterns of brain changes are observed. These include a high proportion of amyloid deposits, medial temporal and parietal brain atrophy, and temporo-parietal FDG-PET hypometabolism, all of which are predictive of short-term conversion to AD. Although imaging modalities show independent contribution towards the prediction of AD, there are considerable redundancies among the neuroimaging modalities for the prediction of AD, and additional costs of combining different imaging acquisitions need to be weighed against the actual gain in prediction accuracy. Accessibility to imaging equipment can also be a limiting factor. The [^{11}C]PiB-PET radioligand, which has a relatively short half-life of decay, can probably only be manufactured at major clinical centers where the imaging equipment is located on-site. However, the development of other ^{18}F -labeled tracers of amyloid with properties more amenable for wider distribution could supplement PiB-PET for clinical use. Of note, the FDA has given conditional support for the use of the [^{18}F]AV-45 PET radiotracer, which should significantly enhance the use of amyloid imaging during the clinical and preclinical stages of AD.

Already, many in the research community appear ready to accept biomarker-aided diagnostic criteria for research purposes, such as has been proposed for the revision of the NINCDS-ADRDA criteria, in part because they provide useful hypotheses for future verification. However, in view of the current evidence, further studies concerning the specific approaches to implement these criteria need to be conducted, including the establishment of quantitative criteria for different biomarkers and their extensive validation (by observing conversion to dementia of the AD type and pathological verification). Only after the clinical utility of such biomarker-aided criteria for the prediction of dementia have been firmly established, can such measures be fully adopted for standardized clinical trials and routine clinical use. Finally, almost all results are reported in carefully selected cohorts, not epidemiological community-based populations; therefore, generalizability of these results to the population as a whole needs to be estab-

lished. Such a validation process will probably take many years. This is especially true when one considers that new biomarkers including blood and CSF assays of compounds not currently known (i.e. evolving from exploratory proteomic approaches) and new imaging radiotracers and techniques (such as PET measures of tau or inflammation, and high-field MRI methods to detect brain amyloid) will probably emerge. Nevertheless, such efforts to develop effective, reliable and robust biomarkers for AD are critical given the significant number of people who will continue to be diagnosed with AD dementia in the coming decades.

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