Amyloid-β Positron Emission Tomography Imaging Probes: A Critical Review

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Abstract. The rapidly rising prevalence and cost of Alzheimer’s disease in recent decades has made the imaging of amyloid-β deposits the focus of intense research. Several amyloid imaging probes with purported specificity for amyloid-β plaques are currently at various stages of FDA approval. However, a number of factors appear to preclude these probes from clinical utilization. As the available “amyloid specific” positron emission tomography imaging probes have failed to demonstrate diagnostic value and have shown limited utility for monitoring therapeutic interventions in humans, a debate on their significance has emerged. The aim of this review is to identify and discuss critically the scientific issues contributing to the extensive inconsistencies reported in the literature on their purported in vivo amyloid specificity and potential utilization in patients.

Keywords: Amyloid imaging, amyloid ‘specific’ imaging probes, Amyvid, critical review, neuropathologic criteria, PIB, silent medial temporal lobe
INTRODUCTION

The rapidly increasing incidence of Alzheimer’s disease (AD) that has accompanied upward trends in life expectancy across the developed world has imposed serious medical, social, and financial burdens that only stand to grow more onerous in the years to come. It is for this reason that the development of biomarkers—positron emission tomography (PET) imaging in particular—for the understanding, diagnosis, and monitoring of various therapeutic approaches to control the progression of AD is a widely accepted need. A great deal of interest has been given to a class of PET biomarkers targeting the amyloid-β (Aβ) peptide, which has been proposed as the root cause of AD. This “amyloid hypothesis” has been a central assumption of research in this field for decades and has underpinned much of the research on the diagnosis and treatment of AD [1]. However, the recent failures of several anti-amyloid therapies in the late stages of clinical testing have raised questions about the utility of measuring amyloid therapies in the late stages of clinical testing of AD [1]. However, the recent failures of several anti-amyloid therapies in the late stages of clinical testing have raised questions about the utility of measuring amyloid plaque as a biomarker of disease progression in patients with substantial clinical impairment. These developments, when combined with the inconsistent data and unexplained anomalies reported in amyloid-PET imaging studies, call for a critical examination of the modality. This review aims to address some of these vital questions regarding Aβ probes by analyzing documented discrepancies between pathological findings, clinical diagnoses, and amyloid imaging observations and examining the essential attributes of PET probes for tissue-specific molecular imaging.

AMYLOID IMAGING IN THE CLINICAL SETTING

On April 6, 2012, the Food and Drug Administration (FDA) approved the clinical use of the Amyvid™ (Florbetapir F 18 Injection) for the evaluation of patients suspected of having AD. This marks the first time a PET biomarker has been approved since 2003, when FDG, sodium [18F]fluoride, and [13N]ammonia were “grandfathered in” based on literature review. Although the approval of [18F]florbetapir can certainly be considered a regulatory breakthrough from this perspective, it has also raised a number of valid questions and caveats regarding its possible uses and potential pitfalls in the clinical setting. The Alzheimer’s Association, in a statement released after the approval of [18F]florbetapir, expressed concern (http://www.alz.org/news_and_events/approval-of-florbetapir.asp) by calling [18F]florbetapir’s approval a double-edged sword, citing fears that unscrupulous operators would make “unrealistic promises” on the value of the tracer when offering imaging services.

This uneasiness is rooted in the severe limitations imposed by the FDA on [18F]florbetapir, the clinical utilization of which has been strictly limited to “ruling out AD.” The explicit exclusion of the more significant applications of “detecting [the] development of dementia” and “monitoring response to therapies” from the FDA’s list of approved indications signifies that [18F]florbetapir scans cannot be used at this time to establish a positive diagnosis of AD or even suggest the likelihood of onset, nor can they be used to validate anti-amyloid pharmacueticals. These restrictions run counter to claims that “the functionality, sensitivity, and specificity of Aβ-plaque imaging agents,” which have been designed “to detect Aβ pathology . . . before the onset of clinically significant cognitive symptoms,” are sufficient for clinical use [2]. In reference to the Aβ imaging agent (N-methyl-11C)2-(4-methylaminophenyl)-6-hydroxybenzothiazole (Pittsburgh compound B or [11C]PiB), it was stated that the probe is “very suitable in the early diagnosis of AD and can be used for the detection of pathological changes before there is a significant loss in cognitive function.” Such claims have frequently been repeated in the literature, which begs the question of why [18F]florbetapir, the first Aβ imaging probe to be reviewed by the FDA, was at first conditionally endorsed and subsequently approved with such stringent limits on its utilization.

Figure 1 (adapted from Fig. 2 from Fleisher et al. [3]) may give an indication as to why the FDA did not approve the use of [18F]florbetapir as a diagnostic tool for AD. The scatter plot presents scan results performed with the subject population likely to receive [18F]florbetapir scans for their clinical assessment: older adults free of dementia (OHC; n = 82), subjects with mild cognitive impairment (MCI; n = 68), and possible AD patients (PAD; n = 60) (as compared with younger adults (YNC)). As seen from the reported results, the vast majority of the subjects at all levels of cognitive impairment will fall within a range of cortical standardized uptake value ratios (SUVR) between 0.80 and 1.50, with the cerebellum—which generally lacks Aβ aggregates of the type to which [18F]florbetapir is expected to bind—established as the reference tissue. These results from Fleisher et al. [3] indicate that [18F]florbetapir would be of limited clinical value for physicians intending to acquire diagnostic information.
Fig. 1. [18F]Florbetapir mean cortical SUVRs of subjects with probable AD (PAD) and mild cognitive impairment (MCI), as compared to older healthy controls (OHC) and young adults (YNC). The top and bottom red lines demarcate the SUVR range that most subjects fall into, regardless of their clinical diagnosis, while the middle line demarcates the highest SUVR values observed in YNC. Note the significant overlap among the groups (From Fleisher et al. [3]; with permission).

on individual patients. The substantial overlap of SUVRs across elderly subjects at all levels of cognitive function (Fig. 1) implies that an individual patient whose SUVR in [18F]florbetapir PET scan falls between 0.8 and 1.5 cannot be clinically classified within any of the aforementioned groups (OHC, MCI, or PAD) based solely on these results. Furthermore, the data displayed in Fig. 1 also support the view that the lone indication that the FDA approved for [18F]florbetapir—ruling out AD with a negative PET scan—may also have severe limitations, as “amyloid-negative” scans would be found in all three major patient groups.

The possible reasons for this extensive overlap between dementia indications and controls (Fig. 1) can be categorized as either (i) biological or (ii) technical (i.e., methodological) in nature.

Biological

It may be that one of the neuropathological pathognomonic elements of AD, Aβ aggregation, does not correlate well with symptomatology. Once the disease process is underway, it is apparent that there are a number of factors, including the accumulation of amyloid plaques, that govern a patient’s symptoms. However, it is less clear which of these factors are responsible for the initiation of AD [4]. The belief that Aβ commences the cascade of neurodegeneration (i.e., the amyloid hypothesis) [1] has most recently led to the conclusion that soluble Aβ oligomers are the form of amyloid that prove deleterious to neurons [5]. An alternative hypothesis suggests that intracellular toxic products of amyloid-β protein precursor processing create neuronal dysfunction and degeneration [6]. But in either case, a linear relationship between Aβ oligomers, internal toxic products, and extracellular amyloid plaques, which are purportedly detectable by current Aβ imaging methods, would not necessarily follow.

Technical

Another possibility is that amyloid plaque density may be a useful measure for diagnosis, but imaging with [18F]florbetapir and other current “amyloid-specific tracers” may not be able to provide an accurate measure of Aβ.

Aβ IMAGING AND NEUROPATHOLOGICAL CRITERIA FOR AD DIAGNOSIS

The diagnostic utility of PET amyloid imaging, which has been claimed to have outstanding specificity for Aβ, appears closely connected to the validity of the amyloid hypothesis [1]. This hypothesis, which proposes that amyloid in its various forms—namely oligomers, fibrils, and neuritic plaques—are etiologically involved in AD has been widely held in the field for a number of decades, but has recently been challenged. Questions regarding the theoretical basis of PET amyloid imaging [7] were met with strenuous objections; advocates of the modality rejected the notion that the uncertainty of the amyloid hypothesis posed a potentially serious limitation, as the hypothesis was “not highly relevant to the feasibility and validity of Aβ imaging” [2]. However, for the intended indication of predicting the development of AD to be viable, the assumption must be made that Aβ is linked to the disease’s clinical symptoms. The same is true for the development of anti-amyloid therapies that amyloid imaging probes have been designed to validate [8, 9]. This assumption has been the subject of increasing scrutiny, especially in light of the recent failures of prominent anti-amyloid pharmaceuticals and the adverse effects observed with anti-amyloid vaccines [10] and the γ-secretase inhibitor Avagacestat [11, 12]. It has even been suggested on the basis of these developments that
PROPERTIES OF THE Aβ PATHWAY TO DIAGNOSIS

Two fundamental reasons about this imaging modality are not appropriate in this case, for at least two fundamental reasons: a) in vitro assay conditions generally used for binding determinations do not resemble in vivo conditions; and b) these experiments are frequently performed in the presence of solvents (e.g., 10% ethanol) and under conditions that are optimized for neuroaggregate binding and for the chemical properties of the tested imaging probe. These experimental conditions are not optimal for the localization of other possible tissue targets (e.g., enzyme-mediated probe transformations), which would necessitate a different set of conditions for their detection and quantification.

ii) As shown by the parallels in their structural backbones, [18F]florbetapir has a close connection with estradiol and potent estrogen analogs (e.g., diethylstilbestrol) (Table 1). However, in spite of these structural similarities with estrogens, no significant reports exist in the scientific literature describing the possibility of any in vivo interaction of the trans-stilbene [18F]florbetapir—or its congener Aβ imaging probe [18F]florbetaben—with estrogen tissue targets, including estrogen receptors and estrogen-related enzymes. This in vivo interaction would be expected because trans-stilbene derivatives bind to estrogen receptors with high affinities [21] and have been known for many decades to have agonistic and antagonistic estrogen effects in vivo. Due to this possibility, [18F]florbetapir’s in vitro and in vivo interactions with estrogen-related targets in the brain are to be expected, and these tissue interactions are likely to influence the in vivo tissue specificity of [18F]florbetapir and—by extension—limit the ability to accurately quantify Aβ plaque loads using this modality.

iii) It has been reported in the literature that [18F]florbetapir is rapidly metabolized peripherally in rodents: the plasma half-life has been shown to be 6 minutes in mice and 11 minutes in rats. Merely 2 minutes after the intravenous injection of [18F]florbetapir, only 56% of the activity in the brain was found to be attributable to the parent molecule, meaning that metabolites account for the remaining 44% [22]. The pharmacokinetic properties of this compound were established in rat plasma, while the identities of its metabolites found in liver microsomes were established by high performance liquid chromatography and mass spectrometry.

[211x433]PROPERTIES OF THE Aβ PATHWAY TO DIAGNOSIS

The diagnostic value of current amyloid-PET imaging probes is also challenged by the established Consortium to Establish a Registry for Alzheimer’s Disease (CERAD) criteria for a neuropathological diagnosis of AD [National Institutes of Aging-Alzheimer’s Association (NIA-AA)], which entails a complex regional localization of tau and Aβ neuroaggregates rather than the mere presence of a widely-distributed amyloid burden [14, 15].

More specifically, it requires an “ABC” score for AD neuropathologic change that incorporates: A) histopathologic assessments of β-amyloidosis in the medial temporal lobe structures, as described by Thal et al. [16, 17]; B) staging of neurofibrillary tangles (Braak tau pathology stages) [18]; and C) scoring of neuritic plaques (based on semi-quantitative determination of senile plaque densities (sparse, moderate, frequent) in at least five neocortical regions, which must include the middle frontal gyrus, superior and middle temporal gyri, inferior parietal lobule, hippocampus, entorhinal cortex, and amygdala) [19].

In its current state, amyloid-PET imaging can only fulfill criterion (C); the available amyloid imaging agents fail to meet criteria (A) and (B), as they are silent in the medial temporal lobe and insensitive to tau deposition [2]. For this reason alone, current Aβ imaging technology is not suitable as a stand-alone method of diagnosis for AD. Although this caveat has been stated repeatedly in the literature, there is a common misconception—often seen in the portrayal of these probes in the media—that amyloid imaging offers a path to diagnosis.
The brain penetrability of $^{18}$F-florbetapir’s peripheral metabolites was confirmed independently by investigators of Avid Pharmaceuticals. In this study [23], at least two of the $^{18}$F-florbetapir metabolites identified were shown to have good blood-brain barrier (BBB) penetrability in the mouse brain. In vivo radiodefluorination was also reported to have been observed with both $^{18}$F-florbetapir and the closely related probe $^{18}$F-florbetaben [24].

Considering the labile nature of pegylated-O-substitution and the reactivity of trans-stilbenes with metabolic enzymes, the possibility that stilbene molecules such as $^{18}$F-florbetapir and $^{18}$F-florbetaben are metabolized in vivo is not altogether surprising. However, the implications of metabolite activity in the brain being as high as 44% only 2 minutes after the injection of $^{18}$F-florbetapir are stark and unsettling. Considering the high peripheral metabolic rate reported in rodents and the high BBB penetrability of these metabolites, it is reasonable to assume that the percent brain accumulation of metabolites would increase to even greater levels over time. At 40 to 60 minutes—the time frame when Logan analysis for quantification (e.g., distribution volume ratio (DVR)) is made [25]—only a small percentage of the activity observed in all regions of the brain PET scan would be from the parent molecules (i.e., $^{18}$F-florbetapir or $^{18}$F-florbetaben), effectively eliminating any possibility of accurately estimating $A_42$ loads.

Limited BBB transport of metabolites would be acceptable as long as their transport is accounted for in a tracer kinetic model for quantification, but the magnitude of the reported peripheral metabolism for $^{18}$F-florbetapir and $^{18}$F-florbetaben makes it highly unlikely that this would be the case. It is to be expected that metabolism translation from rodents to primates would favor higher probe stability in humans, but the quantitative nature of this critical information has not been presented.

### Table 1

<table>
<thead>
<tr>
<th>Molecular Structures of Amyloid Imaging Probes</th>
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<tr>
<td><strong>Benzothiazoles and Related Heterocycles</strong></td>
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<tr>
<td><img src="image1.png" alt="Chemical Structure 1" /></td>
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<tr>
<td><img src="image2.png" alt="Chemical Structure 2" /></td>
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<tr>
<td><strong>Stilbenes</strong></td>
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<td><img src="image3.png" alt="Chemical Structure 3" /></td>
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<td><img src="image4.png" alt="Chemical Structure 4" /></td>
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</table>
The rapidly increasing concentrations of brain metabolites of \([^{18}F]\)florbetapir and \([^{18}F]\)florbetaben, which produce high background signal and hinder signal specificity for A\(\beta\) aggregates, could be one of the reasons leading to the difficulties in reading \([^{18}F]\)florbetapir scans. It is therefore a likely contributor to the inter-reader variability [26] that necessitated the development of a special training program and stimulated the commercialization of specially designed software to facilitate brain scan reading with this amyloid probe. For these reasons, a discussion of the significance and possible implications of the presence of these metabolites on the quantification of A\(\beta\) plaque loads is crucial, especially since transport of peripheral metabolites to the brain is one of the most basic considerations in the design and utilization of optimal PET imaging probes [27].

A SECOND FAMILY OF A\(\beta\) IMAGING PROBES: UNCHARGED DERIVATIVES OF THIOFLAVIN-T (THT)

Table 1 illustrates another family of A\(\beta\) imaging probes in addition to the trans-stilbene derivatives represented by \([^{18}F]\)florbetapir and \([^{18}F]\)florbetaben: the uncharged derivatives of the histological dye Thioflavin-T (THT) [28]. A comparison of these two families of compounds as A\(\beta\) imaging probes can be made using an abundance of peer-reviewed studies in the literature. \([^{11}C]\)PiB, which is an uncharged thioflavin analog, was the first reported A\(\beta\)-specific imaging probe—that is, the first probe to bind exclusively to A\(\beta\) aggregates instead of tau or prion fibrils, among other \(\beta\)-sheet-containing amyloid aggregates [29]. Even though the binding of \([^{11}C]\)PiB and other uncharged 6-hydroxy-derivatives of thioflavins (e.g., \([^{18}F]\)futerometamol) was reported to be A\(\beta\)-specific (Table 1), the charged THT dye that they are derived from is known to target \(\beta\)-sheet conformations present in multiple neuroaggregates (e.g., A\(\beta\), tau, prions, etc.). No specific molecular data exist in the literature, however, to suggest that the binding properties of these uncharged 6-hydroxybenzothiazole derivatives (i.e., \([^{11}C]\)PiB and related molecules) would differ from those of THT. Furthermore, no explanation can be found in the literature as to why they would be specific for A\(\beta\) fibrils or to which fibril site they would bind that is not present in the other neuroaggregates.

The claim of specificity on the part of these probes was mainly based on \textit{in vitro} determinations of tissue specimens, as no evidence of \textit{in vivo} specificity is available [2]. Trans-stilbenes derivatives such as \([^{18}F]\)flutemetamol were consequently presumed to be A\(\beta\)-specific probes as well, since they shared similar brain cortical profiles on their PET scans. However, independent evidence in support of this claim is also lacking in the literature.

A key difference between the two families of imaging probes is that the tissue binding of the parent molecules of thioflavin derivatives has been studied more extensively than those of the stilbene derivatives. Studies performed on the former group have suggested that A\(\beta\) binding of THT is either monomeric, dimeric, or micellar [30]. Obviously, neither dimeric nor micellar binding are likely to be achieved under tracer conditions used in \textit{in vivo} imaging experiments, where the molecular recognition mechanism of these uncharged THT for amyloid fibrils would best be explained through the motor rotor effect [31].

It should also be noted that binding to A\(\beta\) fibrils by multiple dyes and probes has complex characteristics [32]. Several studies have suggested multiple binding sites, and at least three distinct types of binding sites have been identified in A\(\beta\)-40 fibrils [31].

These binding sites were differentiated by kinetic displacement studies when bound to A\(\beta\) fibrils, presumably because each dye binds to a distinct site on the A\(\beta\) aggregates. Recent crystallographic determinations confirm the existence of multiple forms of binding of small molecules to amyloids [33]. At this time, it is not known whether the tracers derived from 6-hydroxybenzothiazole used in \textit{in vivo} studies in humans have the same binding affinities for non-A\(\beta\) tissue targets as THT (e.g., binding to acetylcholinesterases [34] and DNA [35]).

KEY QUESTIONS WITH CURRENT AMYLOID IMAGING PROBES

In a recent editorial, Villemagne et al. [2] acknowledged a host of fundamental limitations involving all “A\(\beta\) specific” imaging probes, including: A) the extensive and uncharacterized uptake of A\(\beta\) PET tracers in the white matter of the brain; B) tissue target specificity of imaging results with the different tertiary forms of A\(\beta\) found at autopsy; and C) the lack of data in support for the \textit{in vivo} specificity of these PET probes.

The extensive and uncharacterized uptake of amyloid probes in the white matter

A pervasive finding in amyloid-PET imaging that appears to affect both dementia patients and controls...
is the substantial retention of these radiotracers in the white matter, a region that is largely spared by Aβ pathology. This phenomenon of apparent non-specific binding in the white matter, which is observed in all existing Aβ-specific PET tracers to varying degrees, is generally accepted and has not been a point of contention [2, 36]. The implications of this recognized non-specificity, however, are a different matter in light of the aforementioned evidence of rapid peripheral metabolism and significant BBB transport of metabolites.

It has been maintained that the unexplained retention in the white matter does not pose a significant threat to the diagnostic accuracy of Aβ PET tracers, and that “this limitation has not proven to be a major hurdle to the quantitation of Aβ deposits in cortical gray matter, neither in the in vivo nor in postmortem cross-validation studies, nor in studies on the predictive value of the Aβ imaging findings, with regard to future cognitive decline” [2]. These claims are not entirely consistent with reported findings in the literature: an earlier paper on [11C]PiB by Klunk et al. acknowledged that Patlak-Gjedde models had difficulties quantifying signals in regions with very low intensity due to “a small but significant net accumulation of tracer in the cerebellum, perhaps because of non-specific retention in the white matter” [29]. The strikingly high SUVs of the white matter of both controls and patients hint at the scope of the non-specific binding of existing Aβ tracers, which has also been thought to “limit the sensitivity of PET imaging especially in a prodromal phase of AD when plaque levels might be low” [29].

Furthermore, it has been established that this white matter PET signal produces a significant spill-over effect. Spill-over is an especially crucial concern in AD, since cortical atrophy would intensify the partial-volume effect. The relatively narrow width of the cortical grey matter in AD subjects makes it particularly susceptible to the partial-volume effect and the associated spill-over from the white matter, which demonstrates a level of retention that is at least equivalent to—and in many cases substantially greater than—that of the grey matter [37]. The importance of partial volume corrections in amyloid-PET imaging and the potential for biases that could lead to erroneous inferences about uptake in certain regions has been recently described in the literature [38]. Thus, the dismissal of the potentially disruptive effects of the strong PET signal consistently detected in the white matter does not appear justified.

A common explanation for the high white matter signal proposes that the phenomenon is “largely attributable to slower . . . white matter kinetics” [2, 36]. Blood flow has indeed been shown to be slower in white matter than in grey matter; multiple studies have measured the difference in arterial transit times to be approximately 650 milliseconds [39, 40]. However, the supposition that this disparity in clearance rate is largely responsible for the non-specific signal in the white matter is challenged by a number of findings. Chief among these is the reported difference in white matter retention that is seen in AD patients and controls. A recent [11C]PiB study involving dynamic imaging revealed that the non-specific retention appears appreciably higher in controls at all-time points [40]. Similarly, it has been demonstrated that [18F]florbetaben has significantly higher (p<0.0001) retention in the cerebral and cerebellar white matter of controls [41]. Studies show that cerebral blood flow in the subcortical regions of AD patients is either lower than or equivalent to what is observed in healthy controls, so it would be expected that non-specific retention would either be equal in the two groups or higher in patients [36, 41]. The observation of higher non-specific retention in controls therefore appears to be inconsistent with the hypothesis that a slower clearance rate is primarily responsible.

The implication that extensive non-specificity underlies the hitherto unexplained retention in the white matter is further underscored by a paper published by Grimmer et al. [42]. The study, which examined white matter hyperintensities on the MRIs of AD patients, concluded that this phenomenon is “a consequence of thickening and sclerosis of arterial small vessel walls . . . which might lead to impaired perivascular drainage of molecules including Aβ”. However, the data revealed no significant association between white matter hyperintensities and [11C]PiB uptake at cross-sectional analysis, indicating that reduced clearance may not be at the root of this anomaly.

By most indications, the extent of the non-specific retention of Aβ imaging agents, particularly those labeled with [18F]fluorine, appears to be more serious than what has been suggested. Although it has been claimed that “the typical retention of Aβ radiotracers in grey matter is . . . at least comparable to threefold higher [than in white matter] in typical Aβ-positive scans” [2], it is well-documented with [18F]-labeled tracers that this is not the case. In AD patients, the average ratios of the signal in the frontal cortices to that in the white matter for [18F]florbetapir, [18F]florbetaben, and [18F]flutemetamol have been reported to have a mean of 0.87, 1.1, and 0.7, respectively [43–46]. In controls, these ratios tumble even further to 0.8, 0.7,
Table 2
Subject differences in brain regional cortical [18F]florbetaben uptake. SUV ratios (MV ± SD) obtained from the 80–90 min p.i. image time frames of the AD patients and healthy controls (HCs), together with individual SUVRs of the one false-negative according to visual PET image analysis by three blinded readers AD patient and the one false-positive HC. Cerebellar cortex served as reference region. Apart from pons/midbrain (singular region), mean SUVRs were taken from corresponding left and right hemispherical SUVRs. Composite SUVRs for neocortex combine volume-weighted means of frontal, parietal, lateral temporal, anterior and posterior cingulate and occipital cortex SUVRs (From [37]; with permission).

<table>
<thead>
<tr>
<th>Area</th>
<th>AD patients</th>
<th>Healthy controls</th>
<th>Effect size</th>
<th>p</th>
</tr>
</thead>
<tbody>
<tr>
<td>Frontal cortex</td>
<td>1.82 ± 0.32</td>
<td>1.39</td>
<td>1.36 ± 0.18</td>
<td>1.67</td>
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<tr>
<td>Lateral temporal cortex</td>
<td>1.57 ± 0.25</td>
<td>1.2</td>
<td>1.24 ± 0.10</td>
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<tr>
<td>Mesial temporal cortex</td>
<td>1.36 ± 0.10</td>
<td>1.22</td>
<td>1.25 ± 0.10</td>
<td>1.23</td>
</tr>
<tr>
<td>Parietal cortex</td>
<td>1.65 ± 0.26</td>
<td>1.26</td>
<td>1.36 ± 0.12</td>
<td>1.54</td>
</tr>
<tr>
<td>Occipital cortex</td>
<td>1.56 ± 0.16</td>
<td>1.4</td>
<td>1.37 ± 0.09</td>
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</tr>
<tr>
<td>Anterior cingulate cortex</td>
<td>1.82 ± 0.31</td>
<td>1.49</td>
<td>1.43 ± 0.15</td>
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</tr>
<tr>
<td>Posterior cingulate cortex</td>
<td>1.89 ± 0.45</td>
<td>1.34</td>
<td>1.37 ± 0.13</td>
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</tr>
<tr>
<td>Caudate head</td>
<td>1.55 ± 0.25</td>
<td>1.39</td>
<td>1.18 ± 0.12</td>
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<tr>
<td>Putamen</td>
<td>1.40 ± 0.25</td>
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<tr>
<td>Thalamus</td>
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<td>1.59</td>
<td>1.59 ± 0.12</td>
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<tr>
<td>Pons/midbrain</td>
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<td>2.17</td>
<td>2.05 ± 0.20</td>
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<tr>
<td>White matter (centrum semiovale)</td>
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<td>2.4</td>
<td>2.18 ± 0.25</td>
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<tr>
<td>Neocortex</td>
<td>1.72 ± 0.28</td>
<td>1.33</td>
<td>1.35 ± 0.12</td>
<td>1.53</td>
</tr>
</tbody>
</table>

and 0.4, indicating that tracer retention in the white matter is in fact markedly higher than that in the grey matter in all cases. These findings are in line with other [18F]florbetaben and [18F]flutemetamol studies that reported that the retention in the white matter far outstripped any cortical region in both AD patients and controls [37, 45].

As illustrated in Table 2 from a [18F]florbetaben study performed by Barthel et al., tracer retention in the white matter has been observed to be significantly higher than that in the frontal, temporal, and parietal lobes in Aβ-positive cases [37]. Similar SUVR values for white matter [18F]florbetaben were also more recently reported in a comparative study of [11C]PiB and [18F]florbetaben in AD and controls [47]. Similarly, Vandenberghe et al. [45] (Graph D in Fig. 2) report that accumulation of the fluorinated PiB derivative [18F]flutemetamol is significantly higher in the white matter than in all cortical areas in AD and MCI patients.

This signal is also highly variable, which insinuates that, similar to [18F]florbetapir and [18F]florbetaben, the imaging probe [18F]flutemetamol is subject to a more complex mechanism of accumulation than has been suggested. The range of SUVRs for [18F]flutemetamol in the white matter is shown to lie between 2.0 and 3.5, as compared with the range of 1.0 to 3.0 for the cortices and striatum. It is also illustrated that non-specific signals in the white matter are substantially more intense with [18F]flutemetamol than with [11C]PiB [44]. These findings are inconsistent with the expectation for the simple non-specific binding that is typically characterized by multiple tissue binding sites with low binding affinities, which produce an essentially uniform distribution of activity in the cortices, subcortical area, and white matter (i.e., SUVRs approaching 1 throughout the brain). This large white matter signal would unavoidably cause spill-over and partial volume effects over any poten-
cial cortical signal, making any reading determination unreliable and inconsistent among readers.

The white matter binding pattern observed with these amyloid probes points to high-affinity targets other than Aβ aggregates. Although it has been hypothesized that “thiourated ligands [are] more prone to [having] high non-specific white matter binding” due to their higher lipophilicity, no evidence has been published to confidently identify the origin of this apparent non-specific retention. There are other accounts of the sources of non-specificity in the white matter, including a recent paper that attributed [11C]PiB white matter retention to myelin [48]. Similarly, other studies with 4,4′-disubstituted stilbenes [49, 50] provide direct evidence in vivo and in vitro that these derivatives do bind to myelin with high affinity (Kᵢ in the nM range). These findings provide additional evidence that specific non-Aβ tissue targets rather than the non-specific binding are responsible for the high retention of Aβ imaging probes in white matter.

Tissue target specificity of imaging results with the different tertiary forms of Aβ found at autopsy

The question of the specificity of amyloid imaging probes can be broken down into two components: (1) the in vivo tissue targets of Aβ probes, and (2) the quantitative correlation of regional imaging signals and autopsy findings.

The in vivo binding properties of Aβ imaging probes

The literature contains highly divergent reports of the binding selectivity of the tertiary forms of Aβ for amyloid tracers. Most of these correlational measures have been made with [11C]PiB, but recent correlational studies have also been reported with [18F]Flutemetamol PET [51] or [18F]Florbetapir PET and pathology load [52].

Though it is often claimed that [11C]PiB “binds strongly to fibrillar Aβ” . . . but only weakly to diffuse Aβ plaques” [53], a number of studies have observed significant [11C]PiB retention in correlation with the presence of both fibrillar and diffuse Aβ plaques, leading authors to assert that this tracer “cannot differentiate neuritic from diffuse amyloid plaques” [54]. An affinity for diffuse plaques was suggested in a paper by Kantarci et al. [55], which reported that “diffuse plaques contribute significantly to [11C]PiB signal in Lewy body dementia.”

Similarly, while some studies have suggested that cerebral amyloid angiopathy (CAA) is not detectable by PET using [11C]PiB, others have found “prominent [11C]PiB staining of CAA [56, 57]. Such findings have led authors to conclude that [11C]PiB is not specific for classical plaques” due to the fact that “it additionally binds a range of Aβ containing lesions including diffuse plaques and CAA” [58, 59]. A paper by Sojkova et al. [60] reported that “in a group of 6 individuals with [11C]PiB in vivo imaging and post mortem evaluation, in vivo amyloid imaging showed limited overall agreement with CERAD neuritic plaque ratings. Moreover, the affinity of amyloid tracers for the Aβ42 and Aβ40 isoforms is the subject of some disagreement in the literature. Although it is commonly stated that [11C]PiB labels “both Aβ12- and Aβ140-immunoreactive plaques,” [61] a recent study correlating distribution volume ratios to histopathology found that this was only true in amyloid-positive scans; in amyloid-negative cases, [11C]PiB was only correlated with Aβ42 [62].

A significant number of cases have also been reported where [11C]PiB PET cortical signal is either low or missing (i.e., negative [11C]PiB scans) in the presence of substantial brain Aβ loads at autopsy [8]. These cases generally involve subjects with specific mutations (e.g., E693G substitution in the APP gene—“Arctic mutation” or APPcsc; “Osaka mutation” E693), leading Ringman et al. [63] to reiterate the hypothesis that “different ‘conformations’ of Aβ deposits affect the binding patterns of tracers and that Aβ imaging may not recognize all types of Aβ deposits with equal sensitivity.” They cautioned that “interpreting a negative result is a perilous endeavor” [63]. Similar negative or low binding results have been reported with [11C]PiB scans in transgenic mice with large cortical amyloid deposition [64] and with [11H]PiB in brain specimens of non-human primates [65].

Thus, readers of these amyloid PET scans are left without a clear understanding of what the cortical signal is attributable to. It is uncertain how amyloid probes such as [11C]PiB interact with many forms of Aβ: diffuse and fibrillar, Aβ42 and Aβ40. A possible explanation of these striking contradictions is that the observations and inferences were mostly made by correlating PET imaging results of Aβ tracers with autopsy findings using immunohistochemistry and related techniques. An inherent pitfall of these inconsistent results is that correlative measures do not indicate causality; that is, the inconsistent results may be explained by PET imaging signals not necessarily caused by these aggregates in any of their forms, but rather by tissue targets partially related (e.g., inflammatory tissue targets) or unrelated to these pathologies.
Quantitative correlation of regional imaging signals and autopsy findings

Correlation studies of imaging results and pathology deposition have been termed “semiquantitative” [2]. Therefore, as a part of testing the hypothesis that these probes bind to Aβ, it is necessary to examine and correlate PET signals with pathology distribution on qualitative (i.e., matching spatial distribution of both signals) as well as quantitative levels (i.e., matching densities of both signals). Furthermore, if the hypothesis is correct, then PET imaging should predict the distribution and quantitative levels of pathology post-mortem.

To date, at least one biopsy study [66] and several autopsy studies have reported the correlation of pathology with [11C]PiB in subjects with AD [52, 54, 67], dementia with Lewy bodies [55, 62], Parkinson’s disease with dementia [68], and sporadic Creutzfeldt-Jakob disease [69]. However, with the exception of the study by Ikonomovic et al. [53], which was limited to one AD brain slice, none of these studies mapped out Aβ pathologies in sufficient detail to allow for quantitative regional cortical correlation of amyloid imaging signal with underlying pathology throughout the whole brain, irrespective of the neurodegenerative disease.

Considering the established NIA-AA pathology criteria requirements for the diagnosis of AD [14–19], the question of why Aβ imaging probes have low amyloid binding in the medial temporal lobe is of particular interest. It has been suggested that factors such as cortical atrophy “may contribute to the underestimation of [11C]PiB accumulation in medial temporal lobe” [70]. Other possible factors may include topographical differences in the distribution of Aβ, which is expressed in lower densities in limbic areas such as the hippocampus and subiculum, where neurofibrillary pathology is more prevalent [71]. Yet, in other medial temporal lobe areas, such as the entorhinal, perirhinal, and rhinal cortices, as well as the parahippocampal gyrus, Aβ plaque densities (both diffuse and neuritic) reach levels observed in the lateral temporal lobe and other cortical areas [16, 72]. Partial volume effects are expected in all cortical areas, particularly in severe dementia, but it is considered unlikely that the negligible binding of [11C]PiB in the medial temporal lobe at all levels of cognitive impairment is entirely due to the combined effects of low Aβ plaque densities and atrophy [70].

The typical [11C]PiB pattern in AD, for example, manifests levels of signal that are high in most cortical areas, except in the isocortical medial temporal lobe, an area high in Aβ. These results conflict with data from neuropathological AD studies in the literature (Fig. 3) showing high densities of Aβ plaques in the inferior temporal cortex (Brodmann area 20) and peripheral cortex (isocortical medial temporal lobe, Brodmann areas 35/36) exceeding levels observed in the frontal lobe [71]. Instead, [11C]PiB shows high signal in the precuneus, which has been found to have an Aβ load that is similar to other surrounding neocortical regions [71]. Arnold et al. [71] report that densities of neuritic plaques (defined as Thioflavin-S positive plaques) in both areas of the temporal lobe in AD subjects are at comparable or higher levels than the frontal lobe or precuneus (Brodmann area 7). Several other published studies also describe comparisons of plaque densities in the frontal lobe (Brodmann area 9) and inferior temporal gyrus (Brodmann area 20) that reveal either no difference between the regions or higher densities in the inferior temporal gyrus [73]. This low profile of [11C]PiB accumulation in the inferior and medial regions of the temporal lobe in comparison to the frontal lobe, parietal lobe, precuneus, and lateral temporal lobe has been widely reported in PET studies on AD. Similar patterns have also been demonstrated with [18F]Flutemetamol, [18F]Florbetapir, and [18F]Florbetaben PET scans.

In the early stages of dementia, including MCI, [11C]PiB cortical accumulation has a bi-modal character that resembles the typical profile of either AD patients (“positive PiB scan”) or controls (“negative PiB scan”) [74]. In these cases, discrepancies are also observed with known neuropathological profiles that are defined by the NIA-AA diagnosis criteria and Thal et al. [14–19]. Figure 3 shows that in Thal Aβ phase 2 (the equivalent of Braak amyloid stage B), medial temporal lobe structures already present accumulation of Aβ deposits while Aβ deposition in neocortex remains limited. Indeed, at this early stage, Aβ deposition is confined to a small number of neocortical brain regions, with many cortical brain regions still free of Aβ [16, 17].

This discrepancy between neuropathological studies and [11C]PiB (and other Aβ imaging probes) binding is most evident in voxel-to-voxel comparisons where [11C]PiB binding in the medial temporal lobe—comprising both limbic (i.e., hippocampus, amygdala, entorhinal cortex) and neocortical (i.e., Brodmann 35/36) areas—fail to separate control and AD groups (Fig. 4) [75, 76]. The inconsistency between imaging results and neuropathological findings raises the crucial question of why [11C]PiB and related Aβ imaging probes produce such weak signal in an area rich in both diffuse and neuritic amyloid...
Fig. 3. I. Phases of β-amyloidosis in medial temporal lobe (MTL) structures as described by Thal et al. [17] are incorporated in the NIA-AA revised neuropathologic criteria for AD diagnosis [14, 15]. II. Relationship between cortical and MTL Aβ deposition is demonstrated on coronal brain tissue slices of a whole hemisphere stained with Campbell-Switzer silver stain (see Thal et al. [16] for details) on an example of Thal Aβ Phase 2 case (left image) and an example of widespread amyloidosis in a Thal Aβ Phase 4 case (right image). Note that in Thal Aβ Phase 2 (equivalent to Braak amyloid stage B), some MTL structures (red arrowheads) already have Aβ deposits whereas Aβ deposition in neocortex is still confined to a limited number of neocortical brain regions (e.g., frontal lobe, temporal lobe) and many cortical brain regions still appear free of Aβ deposits (e.g., cingulate gyrus; large black arrows). In Thal Aβ Phase 4 (equivalent to Braak amyloid stage C), MTL structures beyond hippocampus show dense Aβ deposits comparable in density to other neocortical structures (right image). III. Correlation of Thal phases of β-amyloidosis in MTL structures with Braak NFT stages (from [17]). IV. Correlation of both pathology measures with pre-mortem CDR scores (from [16]). (This Figure contains parts of Figures from [16] and [17], with permission).
Fig. 4. Mismatch between neuropathological results and ‘amyloid specific’ imaging probe signal in the medial temporal lobe. In AD medial temporal lobe structures have abundant Aβ deposits, both diffuse and neuritic plaques, comparable in density to those observed in the cortex as shown for temporal lobe by Price and Morris [72] (upper image in the right column) and Thal and colleagues [16] (lower image in the right column). Left column shows coronal view of the [11C]PiB PET image (upper image), of the MRI image from the same subject (lower image) and overlay of [11C]PiB PET image on MRI imaging for anatomical reference. Images in the central column show expanded temporal lobe structures on overlay of [11C]PiB PET image on MRI image (upper insert) and on the MRI image (lower image). The mismatch between amyloid PET imaging results and pathology in medial temporal lobe is demonstrated on an example of the AD [11C]PiB PET scan (upper image in the left row). The [11C]PiB PET signal is highly elevated throughout the cortex but is missing in the medial temporal lobe. Similar results have been reported for other “amyloid-specific” imaging probes. (This Figure contains parts of Figures from [16] and [72]; with permission).

In vivo specificity of amyloid probes

As pointed out by Villemagne et al. [2], dose-escalation and competitive displacement studies have not been performed on any Aβ imaging agents, meaning that the in vivo binding properties of these tracers have not been fully characterized. In addition to the reported inconsistencies between the signal produced by Aβ imaging probes and tissue localization of pathology, the recent critical finding that the observed [11C]PiB accumulation in stroke [77] and meningioma patients [78] (Fig. 5) is not due to the presence of Aβ in the lesions or tumors further underscores the possible role of tissue targets other than Aβ in causing tracer accumulation in the human brain. These observations add to those discussed earlier for white matter signal [49, 50] and demonstrate that [11C]PiB and other Aβ-specific probe retention, as observed in PET images, is not necessarily indicative of Aβ content. Logically these PET scans require caution in their interpretation. The observation that 6-hydroxybenzothiazoles, the parent molecular structure of [11C]PiB and [18F]flutemetamol, are excellent substrates of brain estrogen sulfotransferase (SULT1E1) offers an interesting alternative interpretation to the retention of these imaging probes in the brain in a manner similar to FDG for hexokinase [79]. Published results in rodents
suggest that [\(^{11}\)C]PiB and its derivatives are retained in vivo in brain tissue mainly as 6-O-sulfates—-with a small percentage of 6-O-glucuronidates—producing a signal unrelated to A\(\beta\) deposition. There is evidence that the sulfation of \([\(^{11}\)C]PiB is not limited to rodents; rather, it appears that the compound is subject to similar peripheral metabolism in humans, producing 6-O-sulfate (or \([\(^{11}\)C]PiB-6-O-sulfate) as the main \([\(^{11}\)C]PiB metabolite in plasma. This stable, charged metabolite of \([\(^{11}\)C]PiB, as well as all other 6-hydroxy congeners, does not cross the BBB. However, the presence of estrogen sulfotransferase has been independently confirmed in the human brain, which implies that 6-O-sulfation of \([\(^{11}\)C]PiB and related 6-hydroxy derivatives\) (Table 1) is also feasible in the brains of human subjects.

It is worth noting that the sulfation of \([\(^{11}\)C]PiB was initially demonstrated in the rat brain, but later dismissed in humans based on a single reported autopsy experiment on a frozen brain tissue specimen\(\) [81]. Due to enzyme inactivation, the brain estrogen sulfotransferase and ATP-dependent sulfation of \([\(^{11}\)C]PiB and its 6-hydroxy analogs generally cannot be measured in tissue specimens unless they are fresh, which is a rarity in human brain autopsies. For this reason, brain specimen preparations that are appropriate for detection of amyloid aggregates (i.e., formalin/sucrose fixed or frozen for long periods of time) would not be conducive to the detection of other tissue targets, such as the active sulfotransferase-mediated trapping of \([\(^{11}\)C]PiB and related 6-hydroxy derivatives.

These observations in rodents, along with the presence of estrogen sulfotransferase in the human brain, add to the need for unequivocal confirmation that the PET signal obtained in vivo in the human brain with \([\(^{11}\)C]PiB or any of its radiofluorinated 6-hydroxy derivatives (e.g., \([\(^{18}\)F]flutemetamol, \([\(^{18}\)F]AZD4694) [79], and \([\(^{11}\)C]AZD2184\) (Table 1) is attributable to binding of the parent molecular imaging probe to brain A\(\beta\) aggregates and not to the action of estrogen sulfotransferase, among other possible tissue targets. This may further explain why all 6-hydroxysubstituted benzothiazole/benzoxazoles in current use have in vivo behavior similar to \([\(^{11}\)C]PiB\) [82].

Using thioflavin (or structurally related) derivatives that lack 6-hydroxy substitution but retain tight binding to A\(\beta\) fibrils would remove estrogen sulfotransferase as a tissue target and help clarify uncertainties as to the origin of the brain imaging signal obtained. This information is already available in published studies: Klunk et al. [83] report that the “binding of 2-(4′-methylaminophenyl)benzothiazole (BTA-1) [a thioflavin analog of PiB lacking the 6-hydroxy group] to AD brain is dominated by a specific interaction with A\(\beta\) amyloid deposits.” BTA-1 has a very tight nanomolar binding (\(K_d\)) to A\(\beta\) fibrils in vitro and to AD brain homogenates, properties that parallel the A\(\beta\) binding parameters for \([\(^{11}\)C]PiB (Fig. 6). The absence of the 6-hydroxy-substitution, present in \([\(^{11}\)C]PiB, means that BTA-1 is not a substrate for estrogen sulfotransferases, making it an appropriate candidate for specific A\(\beta\) binding in vivo. Even though the A\(\beta\) binding properties of BTA-1 and PiB are similar [83], Neumaier et al. [84] report that \([\(^{11}\)C]BTA-1 PET imaging properties are quite different from those of \([\(^{11}\)C]PiB, particularly with regard to cortical distribution and SUVR (Fig. 6). Therefore, since \([\(^{11}\)C]BTA-1 binding to AD brain is dominated by a specific interaction with A\(\beta\) deposits, as reported by Klunk et al. [83], the differential cortical distribution and the much higher cortical accumulation of \([\(^{11}\)C]PiB in the AD brain warrants explanation. Since SUVRs are relative measurements (e.g., cortical signal versus cerebellum), differences in peripheral metabolism are not expected to substantially affect (semi)quantitative interpretation of PET signals. It needs to be investigated whether, as suggested by the rodent studies [79], this disparity is attributable to the sulfation of \([\(^{11}\)C]PiB in the human brain.

CONCLUSIONS

The promotion of research on PET biomarkers for all amyloids (i.e., A\(\beta\), tau, prions, etc.) is of the utmost importance for the progress of the field. A fundamental advantage of PET that distinguishes it from other imaging techniques is its ability to quantify the functional activity of specific tissue targets in human
health and disease. To achieve that goal, there are well-established requirements imposed on the design of molecular imaging probes that have been extensively discussed in the literature [1]. The two most vital requirements for the molecular imaging PET probes designed for the regional visualization and quantification of Aβ in the living human brain are: (a) the in vivo signal obtained with PET is only the result of the probe binding to the Aβ neuroaggregates, and (b) the peripheral metabolism of the probe is minimized or, if it occurs, the metabolites either do not penetrate the BBB or do so in a predictable manner that allows for appropriate quantification of the specific signal emerging from the interaction of the probe with Aβ.

The pertinent question is whether the current Aβ PET imaging probes meet these requirements. The lack of in vivo binding validation of these probes and the consequent deficiency in the understanding of their tissue binding and specificity are genuine concerns. Further complications stemming from the rapid peripheral and central metabolism of these probes and the brain transport of metabolites (e.g., in the case of trans-stilbene probes such as [18F]florbetapir) are severe limitations at the very heart of the tracer design and development. The data collected from in vitro binding determinations and autoradiography findings, which are typically obtained with various concentrations of tracers and solvents not mimicking in vivo conditions, simply cannot supplant the results of necessary in vivo experiments.

There exists a host of apparent contradictions regarding the Aβ specificity of amyloid tracers that require in vivo evidence before the regional accumulation of PET tracer can be confidently interpreted as the result of regional Aβ deposition. Moreover, the absence of an adequate model for transforming regional amyloid tracer accumulation data into a meaningful measure of regional Aβ density has led to the postulation of unsubstantiated inferences about AD biology that seek to explain the observed anomalies. For example, without further neuropathological evidence, the aforementioned overlap among MCI and AD patients and asymptomatic older controls has been interpreted to mean that 30% of normal asymptomatic controls have Aβ brain deposition at levels present in full blown AD [85]. Equally worrisome are the opposite findings where patients with positive evidence of pathology deposition and a clinical AD diagnosis demonstrate negative Aβ scans [62, 63, 66, 67].

The existing NIA-AA criteria for the neuropathological diagnosis of AD confirms the importance of both brain tau and Aβ aggregate deposition, since they...
include (A) β-amyloidosis in the medial temporal lobe; (B) Braak tau pathology stages (I–VI stages); and (C) CERAD neuritic plaque staging [14, 15]. These diagnostic criteria represent a significant challenge to the logic behind the utilization of “Aβ-specific” imaging probes as AD diagnostic tools, which the literature points to as their intended use. Villemagne et al. [2] aptly report, however, that “it may be best to not equate amyloid deposition to clinical diagnosis. Rather than as a method of diagnosis (for AD), it might be best to first think of [11C]PiB retention more fundamentally as a method to detect and quantify brain β-amyloidosis . . . It does not provide a diagnosis by itself.” If diagnosis cannot be performed with Aβ PET imaging probes, it would seem that most of the possible benefits of these imaging agents would be derived from their potential application in validating anti-amyloid drugs, assuming they label brain amyloid deposits in vivo. The promise for amyloid tracers to aid in the development of anti-amyloid drugs has been repeatedly touted in the literature. However, in the most recent FDA AD Drug Development Guidance (http://www.fda.gov/downloads/Drugs/GuidanceComplianceRegulatoryInformation/Guidances/UCM338287.pdf), the agency states that it “will not be in a position to consider an approval (of a therapeutic drug) based on the use of a biomarker as a surrogate outcome measure in AD (at any stage of the illness).”

Furthermore, all the reported clinical trials using anti-Aβ antibody (or related) therapies have fallen short of expectations. The most glaring failures have been the recently reported negative results on the Phase III clinical trials of bapineuzumab and solanezumab [10] and the Phase II evaluation of the γ-secretase inhibitor Avagacestat [11, 12]. Using Aβ PET imaging agents as their standard for quantification, bapineuzumab and other anti-amyloid therapies have been shown to substantially reduce Aβ loads in AD patients while still allowing clinical symptoms to progress.

In summary, without a thorough understanding of the elements to play within a scan, an image cannot offer accurate and reliable information. With regards to amyloid imaging, the in vivo specificity of the probes has not been fully established, the sources of their non-specific uptake have not been identified, and the implications of the physical limitations of standard PET systems have not been considered, all of which limit the clinical utility of the modality. All available evidence indicates that (i) current Aβ PET imaging probes do not fulfill the established neuropathological criteria for successful AD diagnosis [14–18]; (ii) diagnostic use is no longer an intended or plausible application of these probes [2]; (iii) their utilization in monitoring anti-Aβ therapies has been very limited, and (iv) anti-Aβ therapies have been repeatedly shown to be ineffective. Placing the available amyloid-PET probes on the market before these critical issues are resolved would therefore run the risk of eroding the trust of the scientific and medical communities in current and future Aβ probes.

EPilogue

“We absolutely must leave room for doubt or there is no progress and no learning. There is no learning without having to pose a question. And a question requires doubt!” Richard P. Feynman (1918–1988), California Institute of Technology, Nobel Laureate, Physics. From lecture “What is and What Should be the Role of Scientific Culture in Modern Society”, given at the Galileo Symposium in Italy, 1964.

This statement by the renowned scientist Richard Feynman is quite apropos to the current saga of Aβ-specific PET probes. AD and related disorders are critical issues facing the aging population, both in developed and developing countries. However, the available data reported thus far on amyloid imaging probes have generated serious doubts as to their validity and usefulness in their proposed indications relating to AD. These concerns have led to relevant questions that must be addressed to better understand the underlying causes of dementia. Any future progress in the PET imaging of AD and related disorders depends on how well these doubts are cleared and lingering questions are answered.

The diagnosis and treatment monitoring of dementia generally stands to benefit tremendously from PET imaging. However, in light of the discussions presented earlier in this review, it appears likely that current probes will face a host of problems in the clinical setting that will limit its diagnostic and prognostic value. Ultimately, Professor Feynman’s axioms of critically assessing scientific approaches and asking pertinent and valid questions in order to establish scientific truth would be necessary for the development and utilization of a truly successful amyloid-PET biomarker.

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