

Insulin resistance predicts brain amyloid deposition in late middle-aged adults

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Abstract

Background: Insulin resistance (IR) increases Alzheimer's disease (AD) risk. IR is related to greater amyloid burden post-mortem and increased deposition within areas affected by early AD. No studies have examined if IR is associated with an in vivo index of amyloid in the human brain in late middle-aged participants at risk for AD.

Methods: Asymptomatic, late middle-aged adults (N = 186) from the Wisconsin Registry for Alzheimer's Prevention underwent [C-11]Pittsburgh compound B (PiB) positron emission tomography. The cross-sectional design tested the interaction between insulin resistance and glycemic status on PiB distribution volume ratio in three regions of interest (frontal, parietal, and temporal).

Results: In participants with normoglycemia but not hyperglycemia, higher insulin resistance corresponded to higher PiB uptake in frontal and temporal areas, reflecting increased amyloid deposition.

Conclusions: This is the first human study to demonstrate that insulin resistance may contribute to amyloid deposition in brain regions affected by AD.

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Keywords:

Insulin resistance; Amyloid; PiB; Alzheimer's disease; Cognitively normal; Prefrontal

1. Background

The etiopathogenesis of Alzheimer's disease (AD) is partly characterized by extracellular β -amyloid (A β) aggregation and medial temporal lobe atrophy [1]. Insulin resistance (IR) is associated with brain amyloidosis in rodents and humans [2–6]. IR is characterized by the loss of tissue

responsivity to insulin and progressive compensatory peripheral hyperglycemia. Some studies suggest that higher IR is present in AD [1], increases AD risk [7], and is associated with post-mortem A β plaques [8]. IR may increase A β oligomerization and potentiate brain atrophy via neuroinflammation or other downstream effects [9]. Intranasal insulin therapy has been found to conversely increase plasma A β 40/42 ratios and improve cognition [10].

No study has examined the direct association between IR and an in vivo marker of amyloid load in AD-sensitive brain areas in late middle-aged participants. Regions of interest (ROIs) include inferior and medial temporal lobe, ventral prefrontal cortex, and posteromedial cortex [6]. Insulin resistance

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and type 2 diabetes involve multiple mechanistic pathways, and the effect of IR on neural health may differ depending on whether or not hyperglycemia is present [7,11,12].

In this study, we examined if IR was associated with amyloid binding in three AD-sensitive ROIs based on glycemic status, as found in our previous study on IR and glucose uptake [13]. We hypothesized that higher IR, indexed by the homeostatic model assessment of insulin resistance (HOMA-IR) [14], would predict greater amyloid burden using [C-11] Pittsburgh compound B (PiB) [15] positron emission tomography (PET). On an exploratory basis, we also investigated subregions of these areas to provide greater spatial specificity.

2. Methods

2.1. Participants

One hundred and eighty-six late middle-aged adults from the Wisconsin Registry for Alzheimer's Prevention (WRAP) underwent PiB-PET scanning. Demographics are shown in Table 1. Details about selection criteria, recruitment sources, and other aspects are directly discussed elsewhere [16]. Briefly, this ongoing study examines genetic and biological

factors that contribute to the development of dementia-related cognitive decline and neural dysfunction. Participants were classified as either having a positive (FH+) or a negative family history (FH-) of AD. FH+ was defined as having one or both parents with autopsy-confirmed or probable AD as defined by research criteria [17], based on review of medical records and autopsy reports when available. FH- was defined as no formal diagnosis of AD or other significant cognitive decline in either parent, based on information provided by telephone interviews with participants. The inclusion criteria for this study consisted of: normal cognitive function determined by neuropsychological evaluation and consensus meeting similar to Neurological and Communicative Disorders and Stroke-Alzheimer's Disease and Related Disorders Association (NINCDS-ADRDA) recommendations [17], no contraindication for PET or magnetic resonance imaging (MRI) and a subsequently normal MRI scan, no current diagnosis of major psychiatric disease or other major medical conditions (e.g., myocardial infarction, recent history of cancer), and no history of head trauma. The University of Wisconsin Institutional Review Board approved all procedures related to this study. Each participant gave full informed consent before study participation.

2.2. Neuropsychological testing

To confirm that participants in this sample were cognitively normal, the Mini-Mental State Examination (MMSE) and neuropsychological factor scores from the full battery [16] were used (see Table 1). Four cognitive domain factors were derived as described previously [18]: immediate memory, verbal learning and memory, working memory, and speed and flexibility. The individual tests which loaded onto the factors were as follows: (1) Rey Auditory Verbal Learning Test [19], Trials 1 and 2 loaded onto Immediate Memory; (2) Rey Auditory Verbal Learning Test [19], Trials 3 to 5 and Delayed Recall Trial loaded onto Verbal Learning & Memory; (3) Wechsler Adult Intelligence Scale – 3rd edition [20], Digit Span, Arithmetic, and Letter-Numbering Sequencing subtests loaded onto Working Memory; and (4) the interference trial from the Stroop Test [21], and Trail Making Test A and B [22] loaded onto Speed & Flexibility. For the MMSE, a cutoff score of 26 was used based on recommended thresholds from the Alzheimer's Disease Neuroimaging Initiative [23]. It is emphasized here that consensus committee meetings in line with NINCDS-ADRDA recommendations [17] were used to confirm if participants were cognitively normal, rather than just the MMSE and other neuropsychological tests.

2.3. APOE genotype

Apolipoprotein (APOE) genotyping has been described previously [24]. Participants were categorized as "Non-APOE $\epsilon 4$ " (no $\epsilon 4$ alleles) or "APOE $\epsilon 4$ " (at least one $\epsilon 4$ allele).

Table 1

Participant demographics

N	186
Age in y (mean \pm SD)	60.37 \pm 5.63
Gender	
Female	129 (69.4%)
Male	57 (30.6%)
Education (mean \pm SD)	16.61 \pm 2.94
Family history of AD	
Negative	53 (28.5%)
Positive	133 (71.5%)
APOE $\epsilon 4$ status	
Non-APOE $\epsilon 4$	114 (61.3%)
APOE $\epsilon 4$	72 (38.7%)
Diabetes status	
Normoglycemia (<100 mg/dl)	135 (72.6%)
At risk/prediabetes (100–125 mg/dl)	43 (23.1%)
Type 2 diabetes (>125 mg/dl)	8 (4.3%)
DBP	74.11 \pm 8.98
SBP	124.70 \pm 15.19
BMI (mean \pm SD)	28.20 \pm 5.22
Glucose (mg/dl)	95.01 \pm 10.28
Insulin (μ U/ml)	9.53 \pm 7.60
HOMA-IR (mean \pm SD)	2.33 \pm 2.19
Total cholesterol (mg/dl)	202.39 \pm 34.79
MMSE (mean \pm SD)	29.3 \pm 0.96
Speed and flexibility (mean \pm SD)	0.13 \pm 0.86
Working memory (mean \pm SD)	0.20 \pm 1.11
Verbal learning (mean \pm SD)	0.16 \pm 0.95
Immediate memory (mean \pm SD)	0.15 \pm 1.07

Abbreviations: SD, standard deviation; AD, Alzheimer's disease; APOE $\epsilon 4$, apolipoprotein $\epsilon 4$ allele; BMI, body mass index; DBP, diastolic blood pressure; SBP, systolic blood pressure; HOMA-IR, homeostatic model assessment of insulin resistance; MMSE, Mini-Mental State Examination.

NOTE. The four cognitive factors at the bottom of Table 1 are Z-scores. Number of subjects is listed unless otherwise specified in parentheses.

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