African Ancestry Individuals with Higher Educational Attainment Are Resilient to Alzheimer's Disease Measured by pTau181

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Abstract.

Background: Cognitive and functional abilities in individuals with Alzheimer's disease (AD) pathology (ADP) are highly variable. Factors contributing to this variability are not well understood. Previous research indicates that higher educational attainment (EA) correlates with reduced cognitive impairments among those with ADP. While cognitive and functional impairments are correlated, they are distinguishable in their manifestations.

Objective: To investigate whether levels of education are associated with functional impairments among those with ADP. **Methods:** This research involved 410 African American (AA) individuals (Institutional Review Boards 20070307, 01/27/2023) to ascertain whether EA correlates with functional resilience and if this effect varies between *APOE* ɛ4 carriers and non-carriers. Utilizing EA as a cognitive reserve proxy, CDR-FUNC as a functional difficulties measure, and blood pTau181 as an ADP proxy, the non-parametric Mann-Whitney U test assessed the relationship between EA and CDR-FUNC in individuals with advanced pTau181 levels.

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Results: The results showed that EA correlated with functional difficulties in AA individuals with high levels of pTau181, such that individuals with high EA are more likely to have better functional ability compared to those with lower EA (W=730.5, p=0.0007). Additionally, we found that the effect of high EA on functional resilience was stronger in ε 4 non-carriers compared to ε 4 carriers (W=555.5, p=0.022).

Conclusions: This study extends the role of cognitive reserve and EA to functional performance showing that cognitive reserve influences the association between ADP burden and functional difficulties. Interestingly, this protective effect seems less pronounced in carriers of the strong genetic risk allele $\varepsilon 4$.

Keywords: Alzheimer's disease, APOE, biomarker, education

INTRODUCTION

Alzheimer's disease (AD) is the most common type of dementia. Older African ancestry individuals in US having almost two times higher prevalence and incidence compared with older non-Hispanic Whites (NHW), but substantially underrepresented in AD studies [1, 2]. Several factors influence the incidence, prevalence, and risk of AD. These include genetic, clinical, social, behavioral, and environmental determinants [3–8].

The characteristic neuropathological changes of AD, the presence of extracellular amyloid- β deposits and intracellular aggregation of neurofibrillary tangles, can now be measured using biomarkers [9, 10]. These include neuroimaging, cerebrospinal fluid (CSF), and blood-based biomarkers [9, 10]. Recent developments in the field of blood biomarkers showed that blood pTau181 levels strongly correlate with abnormal amyloid accumulation, tau deposition, and neurodegeneration [11–13]. The blood pTau181 levels have been demonstrated to differ significantly between diagnostic groups with amyloid, tau, and FDG-PET scans, further validating pTau181 as a robust biomarker for AD related pathology (ADP) [14, 15].

ADP is typically associated with the decline in cognitive, functional, and behavioral abilities seen in AD [10]. However, some studies have shown that even with advanced ADP, a subgroup of individuals can still maintain reasonable cognition and function, which reflects a better-than-expected performance in relation to the degree of ADP [16, 17]. This inconsistency between ADP and clinical symptoms underscores a possible phenotype characterized by biological pathology without clinical impairment.

'Cognitive reserve' (CR) may account for some of this protection from ADP [18, 19]. CR is thought to be the various thinking abilities that actively compensate for the deficits imposed by the ADP [20]. This compensation yields cognitive performance that exceeds expectations when considering life-courserelated brain changes and brain injuries or disease [21].

Educational attainment (EA) is a commonly used correlate of CR. Higher EA tends to foster resilience against AD clinical manifestations, often resulting in improved cognitive performance compared to those with less education but similar levels of ADP [22-25]. Some studies also suggest an interplay between EA and the APOE $\varepsilon 4$ allele, the strongest genetic risk factor for late-onset AD [26-28]. This complex relationship indicates that CR and genetic factors might interact and influence cognitive outcomes. Intriguingly, research from the MacArthur Study of Successful Aging indicates that the protective effects of education are less pronounced in APOE $\varepsilon 4$ carriers, as measured by the Short Portable Mental Status Questionnaire [29]. Similar findings have been observed in APOE ɛ4 carriers during normal aging, where the protective effects of education are attenuated, possibly correlating with a more rapid cognitive decline [30]. However, the relationship between the APOE $\varepsilon 4$ allele and functional abilities shows mixed results in prior studies. Some research indicates no association between APOE ɛ4 and functional status decline [31, 32], while others report greater difficulties in overall Instrumental Activities of Daily Living (IADL) capacity for APOE ɛ4 carriers [33]. These findings highlight the nuanced interaction between EA and the APOE $\varepsilon 4$ in cognitive performance. However, the impact of EA on specific functional abilities, such as managing finances, engaging in social interactions, and handling personal care, particularly in the context of the APOE ɛ4 allele, remains unexplored. Our study addresses this gap by investigating the novel aspect of how high EA influences these functional abilities in the presence of genetic risk factor APOE. To our knowledge, this specific interplay has not been studied, marking a significant contribution of our research. We specifically investigated whether EA, a proxy for cognitive reserve, promotes

functional resilience in African American (AA) individuals with advanced levels of pTau181 (used as a proxy for ADP) and whether this effect differs between carriers and non-carriers of the *APOE* ε 4 allele.

MATERIALS AND METHODS

Study samples

We ascertained 410 AA participants through AD genetics studies from Wake Forest University (WF, North Carolina), and University of Miami (UM, Florida). These participants represented a diverse AA cohort across both Florida and North Carolina sites. All participants were classified as AA based on selfreported race-ethnicity. It is essential to acknowledge that race-ethnicity is dynamic, shaped by geographic, cultural, and sociopolitical forces and are social constructs. Therefore, while participants self-identified as AA, this classification reflects a complex interplay of social, cultural, and historical influences, rather than a distinct biological identity. The inclusion criteria for the study were having CDR scores, reported years of education, APOE genotypes, and blood pTau181 AD biomarker data. All participants or their consenting proxy provided written informed consent as part of the study protocols approved by the site-specific Institutional Review Boards (20070307, 01/27/2023).

Individual ancestral backgrounds were confirmed using genome-wide genetic data with EIGENSTRAT software [34]. Population substructure data sets were compared with those in the 1000 Genome reference panel YRI (Yoruba from Nigeria) and CEU (Utah Residents with Northern and Western European ancestry) populations. Outliers with respect to CEU population (overlapping within the cluster of CEU) were removed from the datasets [35].

Ascertainment protocols have been consistent across the sites and clinical data assessments capture sociodemographic information including years of education, medical and family history, dementia staging, AD/dementia symptoms, neuropsychological abilities, functional capabilities, and behavioral impairments. The CDR was used to evaluate functional capabilities and served as an outcome measure as described below in the Statistical Analysis section. Biomaterials were also collected at the time of study entry by trained phlebotomists. The detailed description of the study samples, recruitment and cognitive assessment are described elsewhere [27]. 253 participants were diagnosed as non-Cognitively Impaired (nCI), 12 as mild cognitively impaired (MCI), and 145 with Alzheimer disease (AD).

Genotyping

Genome-wide single-nucleotide polymorphism (SNP) genotyping was processed on a Global Screening Array and *APOE* genotyping was performed as described in Saunders et al. [36].

Measurement of serum AD biomarkers pTau181

Serum concentrations of pTau181 were measured using SIMOA chemistry implemented on the Quanterix HD-X instrument (Quanterix, Billerica, MA, USA) [37] according to manufacturer's instructions for the pTau181 Advantage V2 assay (catalog #103714). Samples were randomized according to age, sex, and diagnosis and assayed in duplicate on each plate. Means of two measures were used in analysis. Biomarker levels were log10-transformed to satisfy normality assumptions. Samples were removed if biomarker levels were greater than three standard deviations from the mean of the rest of the samples.

CDR-FUNC

The Clinical Dementia Rating scale (CDR) is a semi-structured interview that assesses cognitive and functional impairment associated with AD to allow for staging based on a global CDR score [38]. The global CDR score is derived from box scores assigned to six domains: Memory, Orientation, Judgment and Problem solving, Home and Hobbies, Community Involvement, and Personal Care. Various studies have examined the utility of these box scores for clinical and research purposes [39-41]. The application of box scores in the assessment of functional impairment has also been studied. For example, Sudo et al. developed a composite score designed to assess functional difficulties (CDR-FUNC) [42]. This score is based on the sum of CDR box scores that are aligned with functional performance (Problem solving, Home and Hobbies, Community Involvement, and Personal Care) and that may have a different trajectory than CDR based cognitive abilities. In addition, Sudo and colleagues reported that the CDR-FUNC score is strongly correlated with the functional performance as measured by the Pfeffer's Functional Activities Questionnaire (FAQ) [42, 43].

Statistical analysis

We hypothesized that individuals with high EA and high levels of pTau181 would present less functional difficulties compared to their counterparts with low EA. We further investigated whether EA correlates with the functional resilience (abilities managing money, interacting with others, and caring for personal needs) differently between *APOE* $\varepsilon 4$ allele carriers and non-carriers. To model our hypothesis, we used EA as a proxy for cognitive reserve, CDR-FUNC as a measure for functional difficulties, and pTau181 as a proxy for ADP. We categorized and quantified the variables as described below:

EA. We stratified years of education into two categories: low EA, individuals up to high school (≤ 8 years), and high EA, individuals high school and beyond it (>8 years) [27].

CDR-FUNC. Using the non-memory components of the CDR, Judgment and Problem solving, Home and Hobbies, Community Involvement, and Personal Care, we formulated a composite score by totaling the individual score of the 4 components with the range of 0 to 3 [42]. The CDR-FUNC score then ranges from 0 to 12, with lower numbers indicating no to mild functional difficulties and higher scores indicating more severe difficulties.

ADP (*pTau181*). We defined advanced ADP individuals as those having log₁₀(*pTau181*) level greater than one standard deviation above the mean. This threshold was selected to identify individuals with notably higher pTau181 levels, reflecting a substantial deviation from the average. This approach is consistent with our aim to categorize those with advanced ADP pathology, while acknowledging the absence of a universally established criterion in ADP research.

Statistical test. To examine the relationship between EA and CDR-FUNC in individuals with advanced ADP, we used the non-parametric Mann Whitney U Test (MW). First, we conducted the analysis using the entire sample of advanced ADP individuals. Next, to determine whether the protective effect of high EA on functional resilience is modified by the presence of the ε 4 allele, a genetic risk factor for AD, we stratified our sample by $\varepsilon 4$ status (carriers and non-carriers) and re-analyzed the association between EA and CDR-FUNC in each subgroup. To better understand the influence of age on our analysis, we stratified our samples into subgroups based on an age threshold of 80. This stratification allowed for a more nuanced exploration of age-specific impacts on the relationship between EA and CDR-FUNC.

Table 1 Study samples characteristics

Characteristics	Low EA	High EA
Count	42	368
Age (mean [SD])	76.7 [6.06]	70.4 [7.3]
Sex (% female)	64.30%	75.00%
APOE ε4 (% ε4 carriers)	28.57%	26.36%
log ₁₀ (pTau181) (mean [SD])	0.20 [0.48]	-0.04 [0.41]
Wake Forest University	30	323
University of Miami	12	45

Table 2 Characteristics of 74 selected individuals with high log10 (pTau181) levels

Characteristics	Low EA	High EA
Count	16	58
Age (mean [SD])	83.6 [7.2]	76.0 [8.2]
Sex (% female)	50.0%	81.0%
APOE ε4 (% ε4 carriers)	62.5%	56.7%
log ₁₀ (pTau181) (mean [SD])	0.69 [0.16]	0.57 [0.17]
Wake Forest University	6	31
University of Miami	10	27

RESULTS

The study sample consisted of 73.9% females, with a mean age of 71.08 (SD = 7.42) years and a mean log10(pTau181) level of -0.01 (SD = 0.42) (Table 1). A total of 74 individuals were classified as having high log₁₀(pTau181) level (mean + 1SD), with 43 carrying the *APOE* ε 4 allele and 31 not carrying it. Our downstream analysis was performed using these 74 individuals, selected based on their classification of having high log10(pTau181) levels (Table 2).

Individuals with high $\log_{10}(pTau181)$ and low EA had significantly higher CDR-FUNC scores (median: 12.0; Q1:8.0 – Q3:12.0) compared to those with high EA (median: 3.75; Q1:0 – Q3:9.0; W=730.5, p=0.0007). The distribution of CDR-FUNC scores between low and high EA in individuals with high $\log_{10}(pTau181)$ is illustrated in Fig. 1A.

To determine the effect of the *APOE* ε 4 allele on the association between EA and CDR-FUNC in high $\log_{10}(pTau181)$ individuals, we examined this relationship separately in ε 4 carriers and noncarriers. We found that in both subgroups, individuals with low EA had significantly higher CDR-FUNC scores (ε 4 allele carriers: median = 12.0; Q1:8.75 – Q3:12.0; ε 4 allele non-carriers: median = 10.75, Q1:8.375–Q3:12) than those with high EA (ε 4 allele carriers: median = 6.0; Q1:3.0–Q3:9.0; W=256.5, p=0.0008; ε 4 allele non-carriers: median = 0,

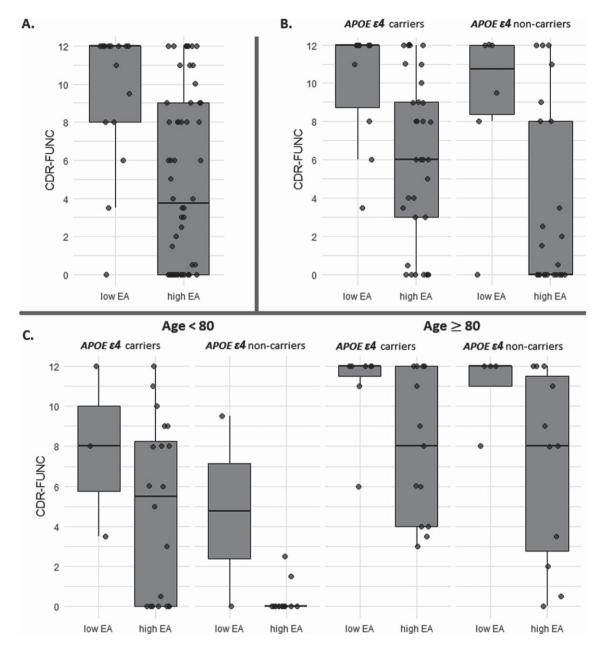


Fig. 1. Box plot for the comparison between educational attainment (EA) levels (low and high) and CDR-FUNC score among advanced Alzheimer disease pathology individuals. (A) compare all individuals with the advanced Alzheimer disease pathology, (B) compare within subgroups stratified by the APOE ε 4 allele, and (C) compare samples further stratified by age to investigate potential age-specific trends.

Q1:0–Q3:8; W = 117, p = 0.029). However, the mean ranks of CDR-FUNC among high EA individuals were significantly higher in $\varepsilon 4$ carriers compared to $\varepsilon 4$ non-carriers (W = 555.5, p = 0.022). We did not observe a significant difference in CDR-FUNC score between $\varepsilon 4$ carriers and non-carriers with low EA (W = 33.5, p = 0.719). Figure 1B shows the distribution of CDR-FUNC scores among advanced ADP individuals with low and high EA, stratified by $\varepsilon 4$ allele.

We further stratified our samples by age to investigate potential age-specific trends. Notably, the patterns distinguishing low and high EA in these age-based subgroups were consistent with those observed in our primary dataset (Age \geq 80: W = 200.5, p = 0.011 and Age < 80: W = 124.5,

p=0.078) (Fig. 1C). Among high EA individuals under the age of 80, significant differences in the mean ranks of CDR-FUNC were found between $\varepsilon 4$ carriers and non-carriers (W=222, p=0.002). However, this distinction was not evident in individuals aged 80 and above (W=79.5, p=0.659).

DISCUSSION

This study found that years of education are associated with better functional abilities in individuals with high \log_{10} (pTau181) levels. This effect was observed in both *APOE* ε 4 carriers and non-carriers, but the protective effect of high EA was stronger in noncarriers. These results extend our understanding of the protective effect that cognitive reserve and EA may have on the functional performance of individuals with AD pathology [16, 17, 22–25]. Based on these findings, it is likely that increased years of education may explain why individuals with known genetic and AD biomarkers on average exhibit better functional performance. It can be inferred that higher education can help individuals maintain their functional abilities even with AD pathology.

Our results highlight a complex relationship among the genetic risk factors, ADP-associated biomarkers, and cognitive reserve and their effect on cognitive and functional outcomes of AD. The study supports the hypothesis, as suggested by Stern et al. [25], that the brain's adaptability and efficiency in cognitive processing can mitigate the impacts of disease-related changes. This is evident in the varying functional outcomes among individuals with high $log_{10}(pTau181)$ levels, where those with high EA demonstrate a more robust cognitive reserve. This reserve, influenced by EA as per the model, is likely to enable the use of alternative neural networks to maintain cognitive function, aligning with the hypothesis by Stern et al. [20]. Particularly, the interaction between EA, genetic factors such as the APOE ɛ4 allele, and AD pathology suggest that cognitive reserve's protective effect is modulated by genetic risk factors. This underscores the role that cognitive reserve, shaped by life experiences and education, plays in determining an individual's resilience to neurological changes and emphasizes its importance in AD management. By investigating potential interactions among EA, ADP, and other genetic and modifiable risk and protective factors for AD, such as lifestyle factors (e.g., diet, exercise, sleep), or environmental exposures (e.g., air pollution, stress) we

can enhance our understanding of the relative significance of these factors in the development of AD. This knowledge can then inform the design of comprehensive interventions aimed at preventing or delaying the onset of AD.

The trend of higher CDR-FUNC scores being associated with lower EA persists across stratified age groups. This suggests that the protective effect of high EA on functional difficulties is a consistent factor, regardless of age. However, it is noteworthy that the sample sizes decrease significantly in the stratified age groups. This reduction in sample size limits the ability to draw significant inferences about the impact of education and APOE e4 carrier status on CDR-FUNC scores within these age brackets. Additionally, CDR-FUNC scores tend to be higher in the older age group (\geq 80 years). This is indicative of a natural progression of cognitive and functional decline with increasing age, which is to be expected in an aging population and particularly in those who are already experiencing AD pathology.

Findings from this study should be interpreted with caution considering several methodological and instrumentation limitations. First, the study population is relatively small and is not representative of all African American populations. Therefore, we are unable to make universal claims that our findings will be observed in all individuals with African ancestry. Additionally, the limited sample sizes in certain subgroups, as indicated in Fig. 1, may constrain the strength of specific statistical inferences. Second, our operationalization of years in school as a proxy for educational attainment, does not capture quality of education, type of school, nature of school curriculum and location of school, all factors that could have impacted the findings. Third, the cross-sectional design of the data precluded us from making any causal claims in the associations between EA and functional status of individual with ADP. Despite these limitations, the strong communitybased representation of our sample, the heterogeneity of participants across multiple sites (Florida and North Carolina) addresses concerns about not having a representative sample, and the use of gold-standard instruments to assess target variables, as well as the public health and clinical value of our findings overshadow these limitations.

In summary, this study shows the importance of using multiple variables such as genetics, education (an established social determinant of health), and AD biomarkers, to have a complete picture of the disease and its cognitive and functional outcomes. Notably, it demonstrates the need for comprehensive approaches to analyze the data, identify patterns associated with a risk of developing AD, and identify the specific combinations of factors most predictive of an individual's risk of developing AD symptoms. Healthcare providers could use this information to identify individuals at high risk of AD and to develop personalized prevention and treatment plans for those individuals.

AUTHOR CONTRIBUTIONS

Farid Rajabli (Conceptualization; Data curation; Formal analysis; Investigation; Methodology; Software; Validation; Visualization; Writing - original draft; Writing - review & editing); Azizi A. Seixas (Conceptualization; Methodology; Supervision; Writing - original draft; Writing - review & editing); Bilcag Akgun (Data curation; Formal analysis; Software; Writing - review & editing); Larry D. Adams (Data curation; Methodology; Writing review & editing); Jovita Inciute (Data curation; Writing - review & editing); Kara L. Hamilton (Data curation; Writing – review & editing); Patrice G. Whithead (Data curation; Writing - review & editing); Ioanna Konidari (Data curation; Writing review & editing); Tianjie Gu (Data curation; Writing - review & editing); Jamie Arvizu (Data curation; Writing – review & editing); Charles G. Golightly (Data curation; Writing - review & editing); Takiyah D. Starks (Data curation; Writing - review & editing); Renee Laux (Data curation; Writing - review & editing); Goldie S. Byrd (Data curation; Funding acquisition; Investigation; Writing - review & editing); Jonathan L. Haines (Methodology; Writing - review & editing); Gary W Beecham (Methodology; Writing - review & editing); Anthony J. Griswold (Data curation; Methodology; Writing review & editing); Jeffery M. Vance (Conceptualization; Methodology; Writing - review & editing); Michael L. Cuccaro (Conceptualization; Investigation; Methodology; Project administration; Writing - original draft; Writing - review & editing); Margaret A. Pericak-Vance (Conceptualization; Funding acquisition; Investigation; Methodology; Project administration; Resources; Supervision; Writing original draft; Writing – review & editing).

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CONFLICT OF INTEREST

The author has no conflict of interest to report.

DATA AVAILABILITY

Data are available through the National Institute on Aging Genetics of Alzheimer's Disease Data Storage Site (NIAGADS) Data Sharing Service (DSS): https://dss.niagads.org/datasets/ng00067/.

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